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(54) **ANTI-ANGIOGENIC PEPTIDES FROM THE N-TERMINUS OF ENDOSTATIN**

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(51) **Int. Cl.**
A61K 38/04 (2006.01)
A61K 38/16 (2006.01)

(52) **U.S. Cl.** **514/2**

(58) **Field of Classification Search** None
See application file for complete search history.

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(57) **ABSTRACT**

Provided herein are anti-angiogenic comprising the N-terminal end of endostatin, nucleic acids encoding the same, pharmaceutical preparations comprising an effective amount of the peptide and nucleic acids and use of the pharmaceuticals in treating or preventing diseases or conditions associated with undesirable angiogenesis.

22 Claims, 8 Drawing Sheets

Figure 1

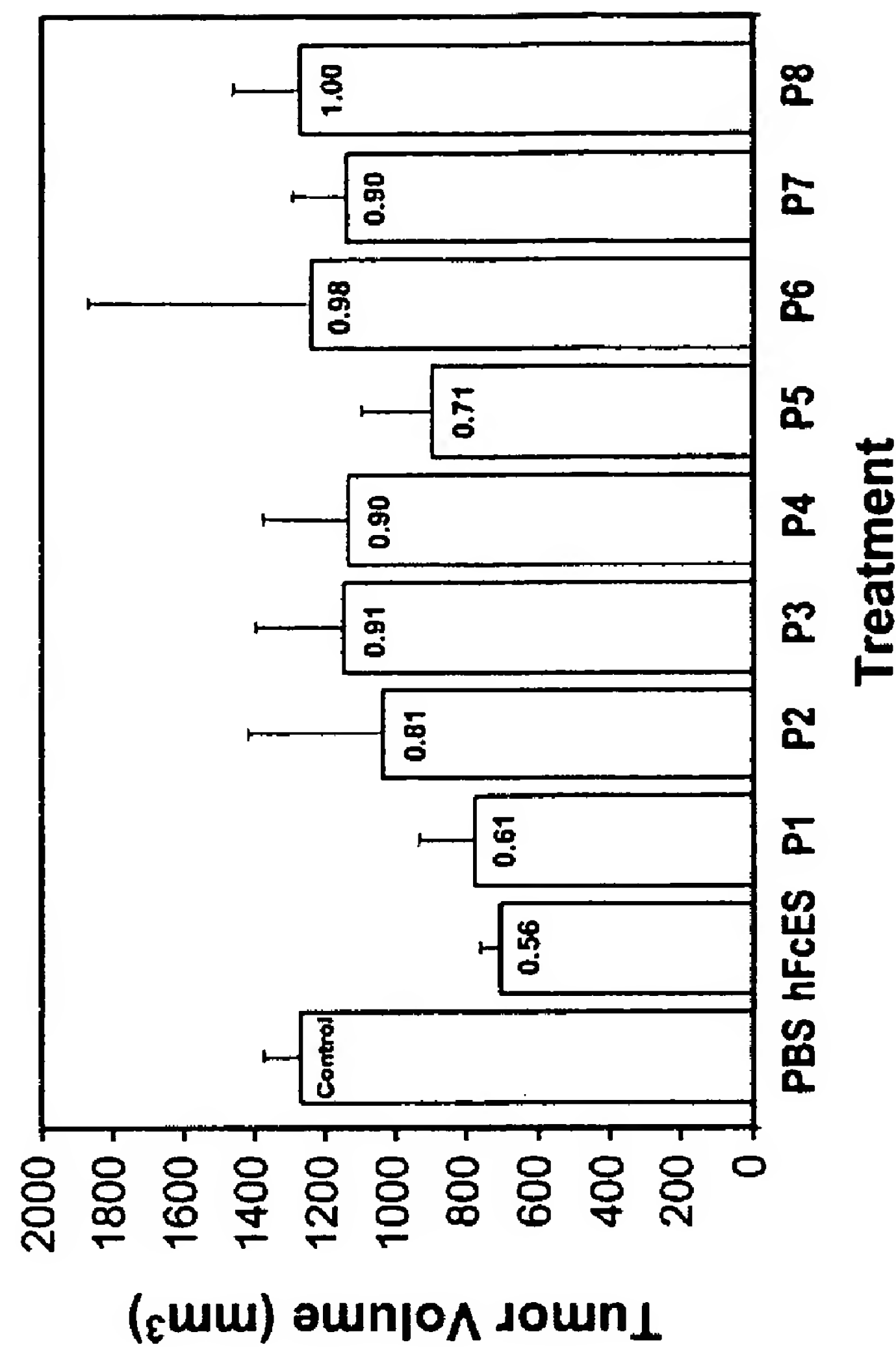


Figure 2

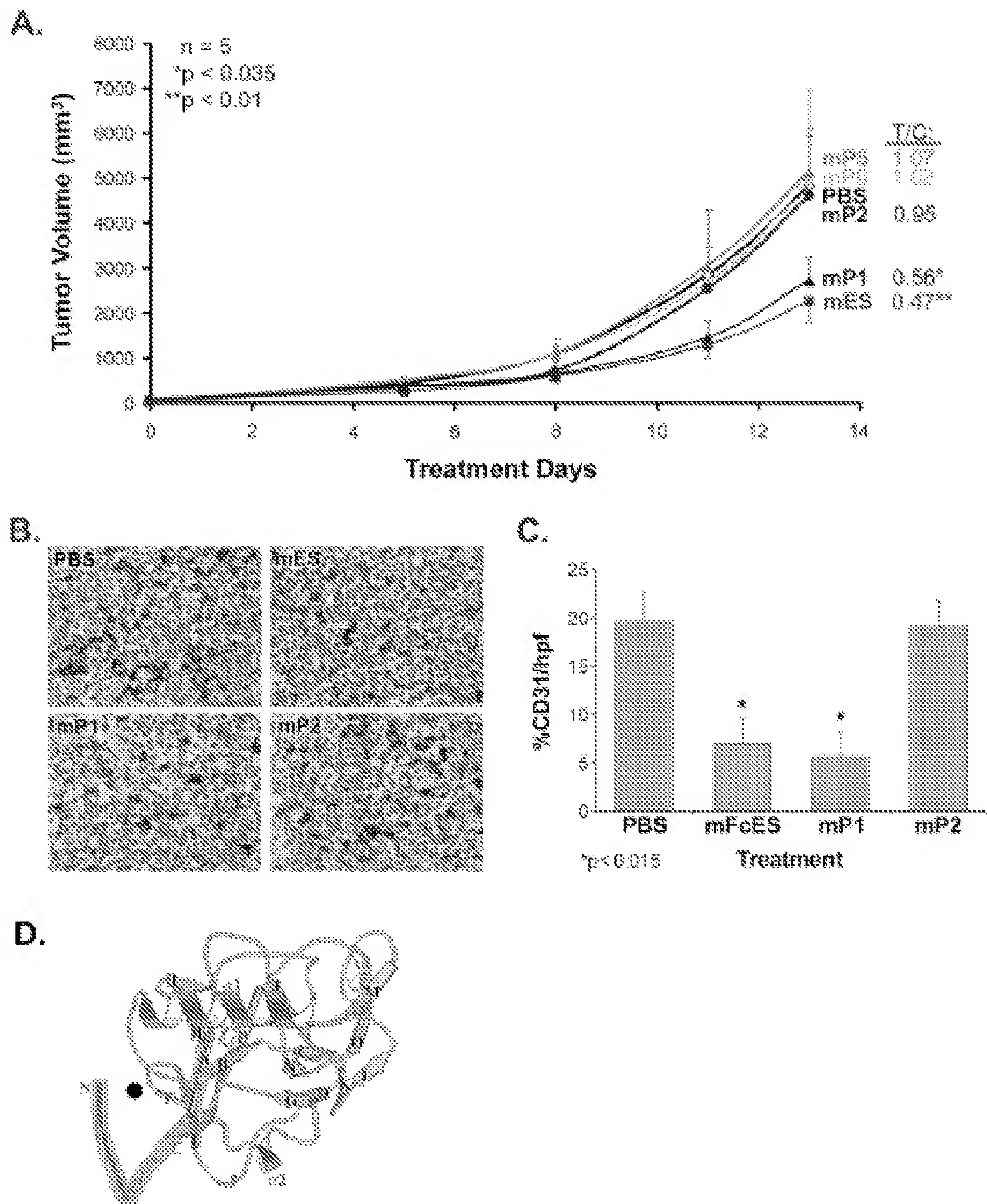


Figure 3

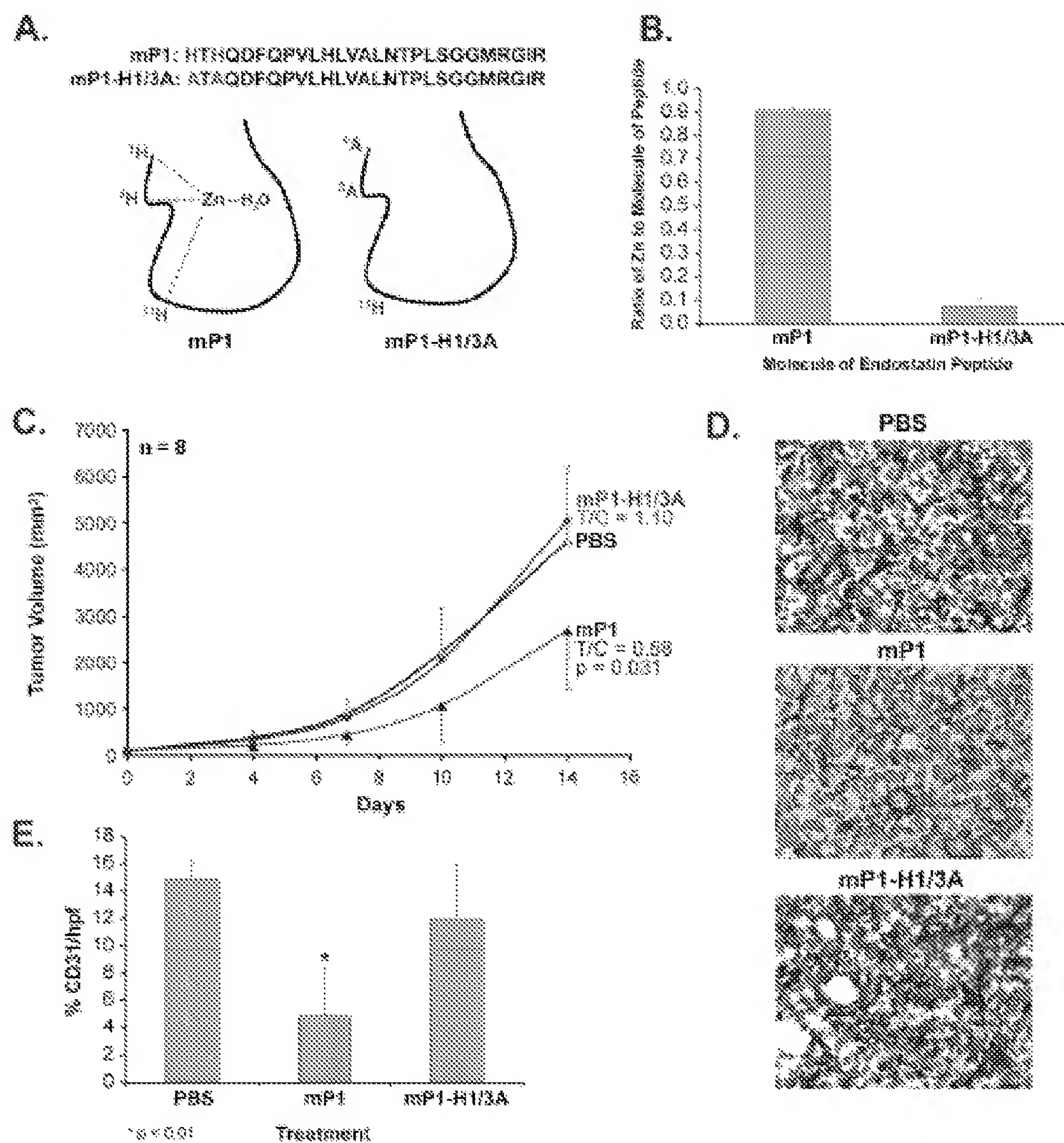


Figure 4

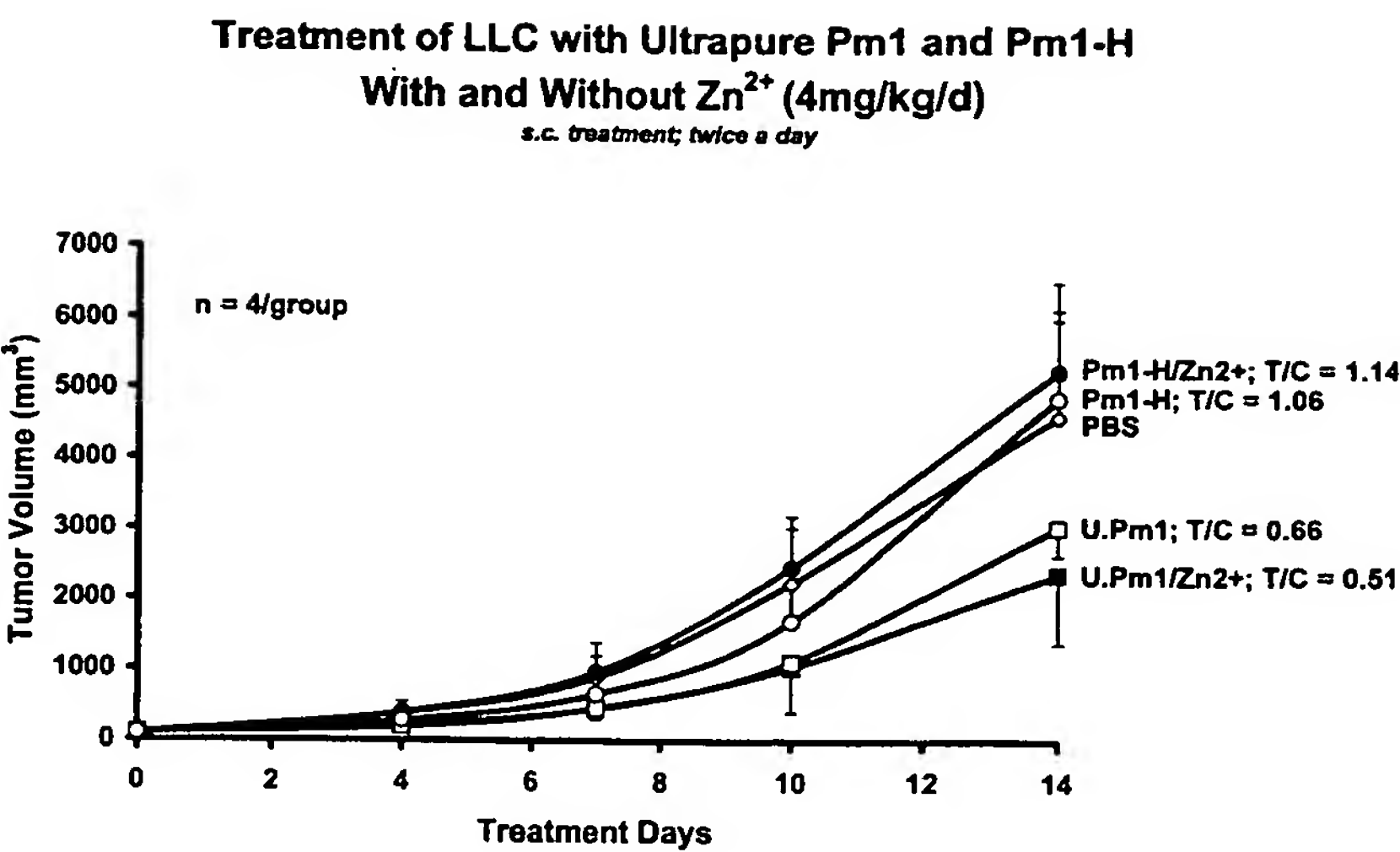


Figure 5

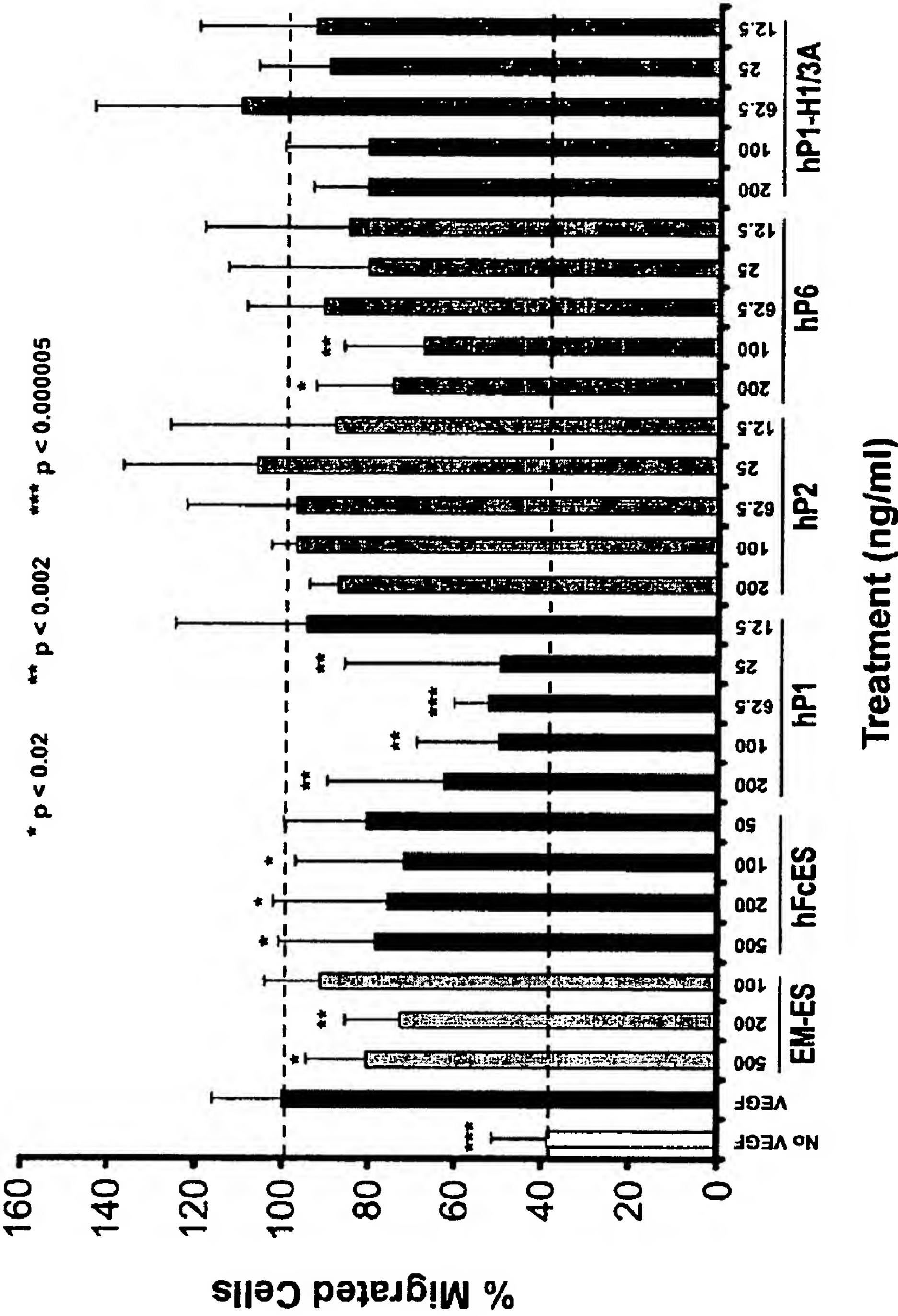


Figure 6

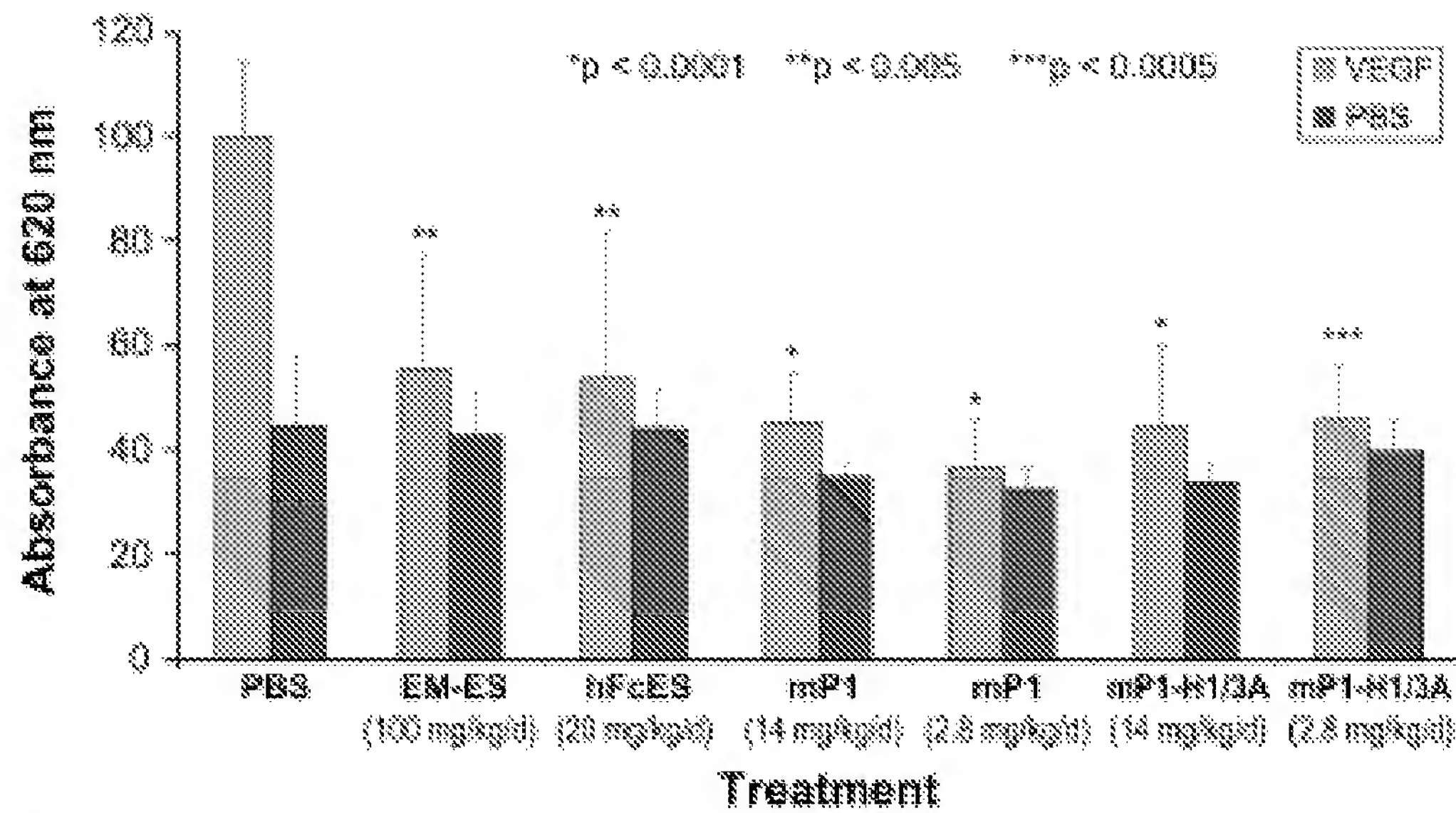
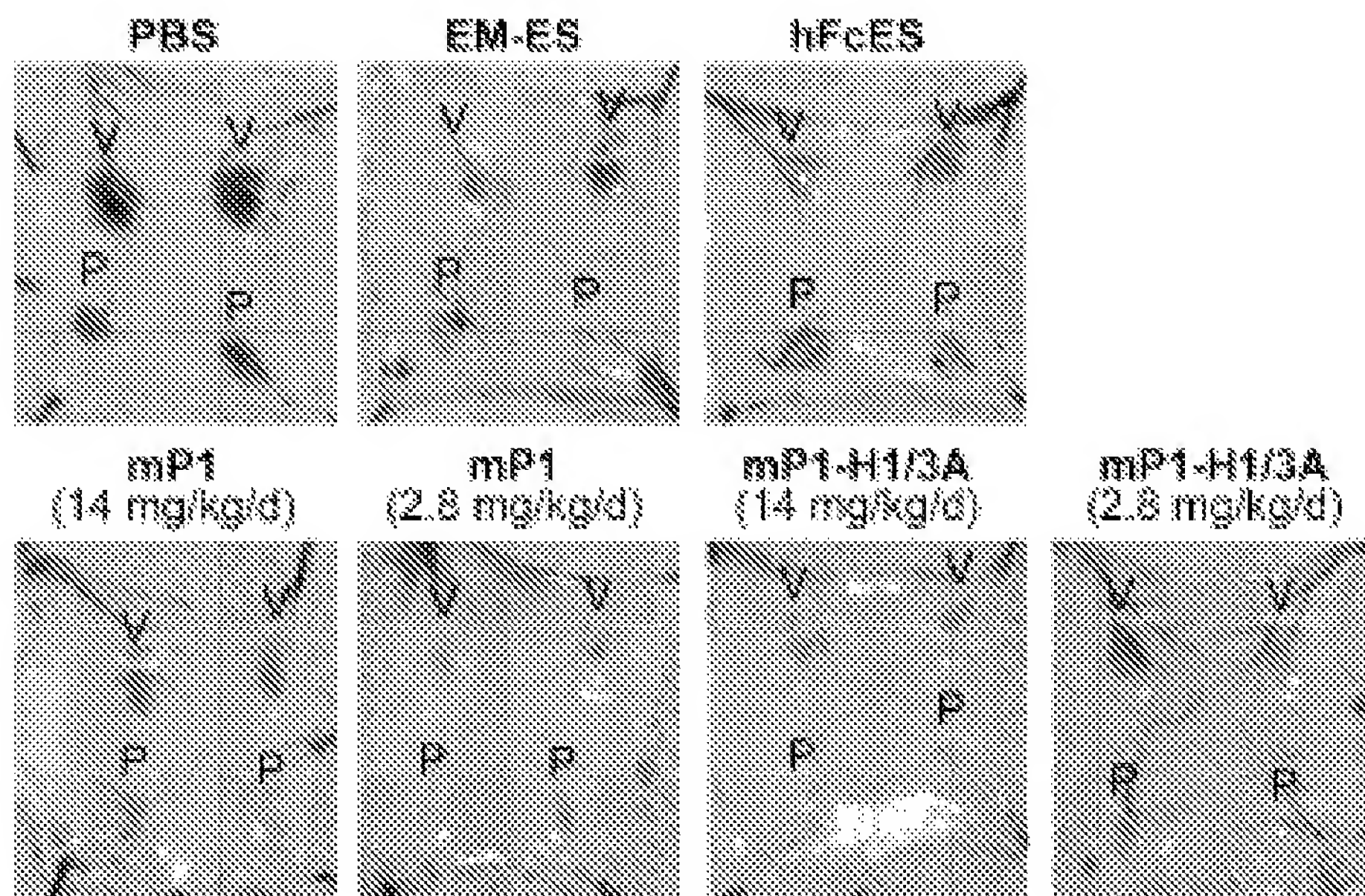
A.**B.**

Figure 7

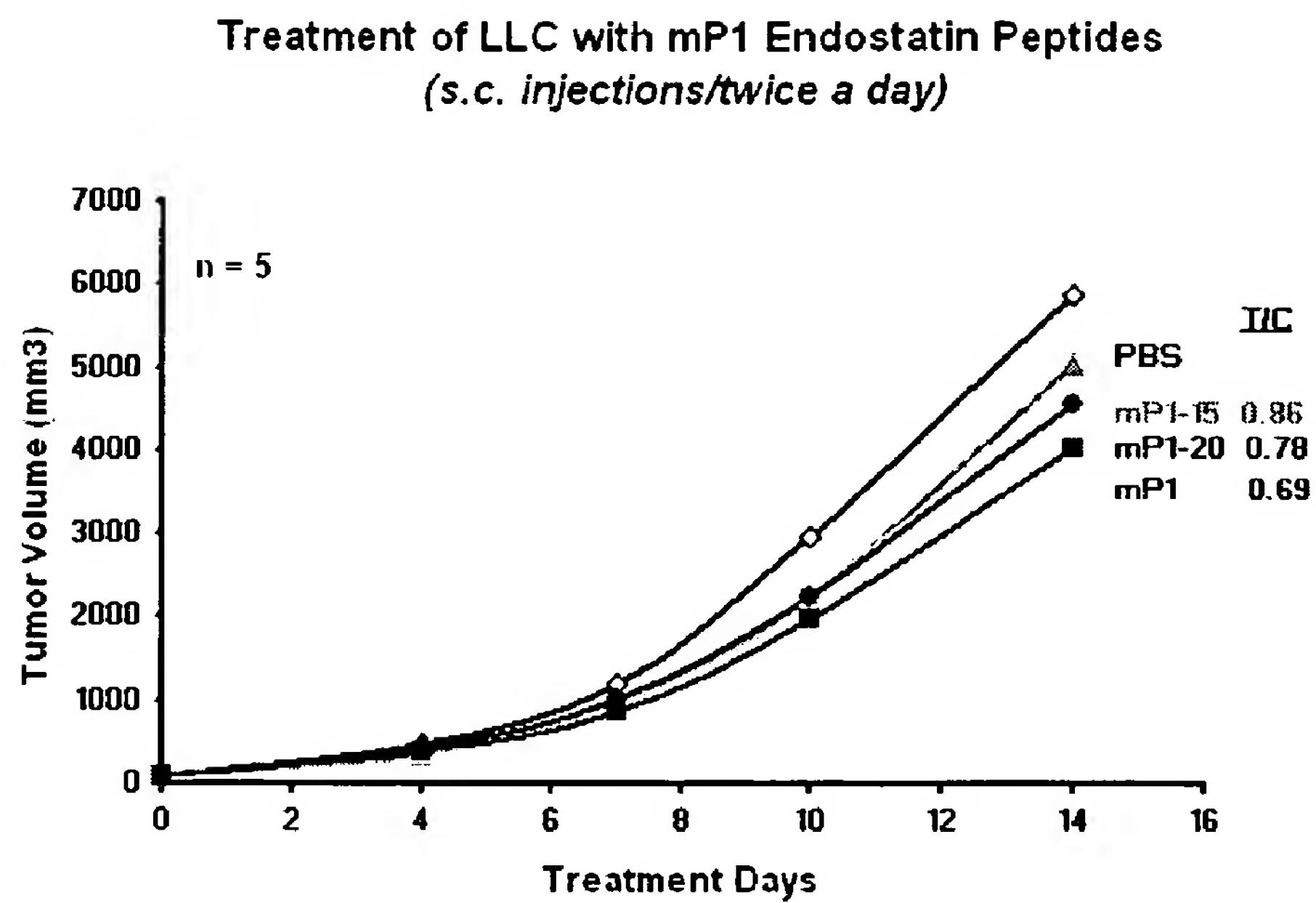


Figure 8

SEQ ID NO: 151 (AF184060 minus first atg):

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58  tcaggcggca tgcggggcat ccgcggggcc gacttccagt gcttccagca ggcgcggggc
118 gtggggctgg cgggcacctt ccgcgccttc ctgtcctcgc gcctgcagga cctgtacagc
178 atcgtgcgcc gtgcgacccg cgcagccgtg cccatcgtea acctcaagga cgagctgctg
238 tttcccagct gggaggctct gttctcagge tctgagggtc cgctgaagcc cggggcacgc
398 atcttctcct ttaacggcaa ggacgtcctg acccacccca cctggcccca gaagagcgtg
358 tggcatggct cggaccccaa cgggcgcagg ctgaccgaga gctactgtga gacgtggcgg
418 acggaggctc cctcggccac gggccaggcc tactcgctgc tggggggcag gctcctgggg
478 cagagtgcgc cgagctgcca tcacgcctac atcgtgctat gcattgagaa cagcttcatg
538 actgcctcca agtag
```

SEQ ID NO: 152 (AAF01310 minus first M):

```
1   hshrdfqpv lhlvalnspl sggmrgirga dfqcfqqara vglagtfrac lssrlqdllys
60  ivrradraav pivnlkdell fpswealfsg segplkpgar ifsfngkdvl thptwpqksv
120 whgsdpngrrr ltesycetwr teapsatgga ysllggrllg qsaaschhay ivlciensfm
180 task
```

SEQ ID NO: 153 (AF257775):

```
1   catactcate aggactttca gccagtgtct cacctgggtg cactgaacac ccccctgtct
61  ggaggcatgc gtggtatccg tggagcagat ttccagtgtt tccagcaagc ccgagccgtg
121 gggctgtcgg gcaccttccg ggcttttctg tcctctagga tgcaggatct ctatagcatc
181 gtgcgccgtg ctgaccgggg gtctgtgccc atcgtcaacc tgaaggacga ggtgctatct
241 cccagctggg actccctgtt ttctggctcc cagggtcaac tgcaaccagg ggcccgcate
301 ttttcttttg acggcagaga tgtcctgaga caccagcct ggccgcagaa gacgtatgg
361 cacggctcgg accccagtgg gcggaggctg atggagagtt actgtgagac atggcgaact
421 gaaactactg gggctacagg tcaggcctcc tcctgtgtgt caggcaggct cctggaacag
481 aaagctgcga gctgccacaa cagctacatc gtccctgtgca ttgagaatag cttcatgacc
541 tctttctcca aa
```

SEQ ID NO: 154 (AAF69009):

```
1   hthqdfqpv lhlvalntpls ggmrgirgad fqcfqqarav glsgtfracfl ssrlqdlysi
61  vrradrgsvp ivnlkdevls pswdslfsgs ggqlqpgari fsfdgrdvlr hpawpqksv
121 hgdspsgrrr mesycetwrt ettgatgqas slsgrlleq kaaschnsyi vlciensfmt
181 sfsk
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**ANTI-ANGIOGENIC PEPTIDES FROM THE
N-TERMINUS OF ENDOSTATIN****CROSS REFERENCE TO RELATED
APPLICATIONS**

This application is a continuation-in-part of PCT/US04/028143, which was filed on Aug. 30, 2004, which claims the benefit of U.S. Provisional Applications 60/499,264, filed on Aug. 29, 2003 and 60/539,213, filed on Jan. 26, 2004; the contents of each application is hereby incorporated by reference in its entirety.

STATEMENT OF RIGHTS

This invention was made with government support under Grant R01 CA064481 awarded by the National Institutes of Health. The government has certain rights in the invention.

BACKGROUND

Endostatin, a 183 amino acid proteolytic cleavage fragment corresponding to the C-terminus of collagen 18, has been the subject of investigation by a number of laboratories because of its anti-tumor activity with no toxic side effects (O'Reilly et al. (1997) *Cell*, 88: 277-285.; Kisker et al. (2001) *Cancer Res*, 61:7669-7674; Dhanabal et al. (1999) *Cancer Res*, 59: 189-197; Yoon et al. (1999) *Cancer Res*, 59: 6251-6256; Folkman and Kalluri, (2003) *Cancer Medicine*, 6th edition, pp. 161-194. Hamilton: B. C. Decker Inc.). A number of anti-angiogenic activities have been reported for this protein, such as inhibition of endothelial cell proliferation, migration, and tube formation. Endostatin also suppresses vascular endothelial growth factor (VEGF)-induced vascular permeability (Takahashi et al. (2003) *Faseb J*, 17: 896-898). However, the mechanism of action of endostatin remains unknown. Endostatin inhibits endothelial cell migration by inhibiting phosphorylation of focal adhesion kinase via binding to $\alpha 5 \beta 1$ integrin (Wickstrom et al. (2002) *Cancer Res*, 62: 5580-5589). It also has been shown that cell surface glypicans are low-affinity endostatin receptors (Karumanchi et al. (2001) *Mol Cell*, 7: 811-822). Endostatin has been implicated in several signaling pathways, such as downregulation of c-myc (Shichiri and Hirata (2001) *Faseb J*, 15: 1044-1053), cyclin-D1 (Hanai et al. (2002) *J Biol Chem*, 277: 16464-16469) and RhoA activity (Wickstrom et al. (2003) *J Biol Chem*, 278: 37895-37901), blockage of VEGF signaling (Hajitou et al. (2002) *Faseb J*, 16: 1802-1804; Kim et al. (2002) *J Biol Chem*, 277: 27872-27879), and inhibition of the wnt-signaling pathway (Hanai et al. (2002) *J Cell Biol*, 158: 529-539). Furthermore, endostatin has been shown to bind and inactivate metalloproteinases (Kim et al. (2000) *Cancer Res*, 60: 5410-5413; Nyberg et al. (2003) *J Biol Chem*, 278: 22404-22411; Lee et al. (2002) *FEBS Lett*, 519: 147-152) and to regulate a spectrum of genes which suppress angiogenesis (Abdollahi et al. (2004) *Mol Cell*, 13: 649-663).

The crystal structures of both murine and human endostatin have been elucidated (Hohenester et al. (1998) *Embo J*, 17: 1656-1664; Ding et al. (1998) *Proc Natl Acad Sci USA*, 95: 10443-10448) and show a noncovalently held dimer at high concentration required for crystallization (Ding et al. (1998) *Proc Natl Acad Sci USA*, 95: 10443-10448). The presence of two disulfide bonds results in a highly folded structure. Endostatin binds one atom of zinc per monomer via the three histidines in the N-terminus of the molecule (histidines 1, 3, and 11) and aspartic 76. The heparin binding property of endostatin is mediated by noncontiguous arginines clustered

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over the three dimensional globular surface of the molecule (Sasaki et al. (1999) *Embo J*, 18: 6240-6248).

Oligomeric endostatin (NC1 and dimer) have been shown to be primarily associated with laminin in the basement membrane (Javaherian et al. (2002) *J Biol Chem*, 277: 45211-45218). This association may be important for some of the biological functions displayed by endostatin. On the other hand, the heparin binding properties of endostatin manifest themselves in its interaction with the cell surface. It is likely that endostatin has a number of biological functions mediated by different regions of the protein.

SUMMARY

The instant disclosure is based on the surprising finding that the N-terminal region of endostatin is responsible for its anti-angiogenic activity. Based on these findings, the disclosure features anti-angiogenic peptides comprising at least about 12 amino acids of SEQ ID Nos. 2 or 4. Other anti-angiogenic peptides comprise at least about 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24 or 25 amino acids of SEQ ID Nos. 2 or 4. Exemplary anti-angiogenic peptides are selected from the group consisting of SEQ ID Nos. 6, 8, 10, 12, 14, 16, 18, 20, 22, 24, 26, 28, 30, 32, 34, 36, 38, 40, 42, 44, 46, 48, 50, 52, 54, 56, 58, 60, 62, 64, 66, 68, 70, 72, 74, 76, 78, 80, 82, 84, 86, 88, 90, 92, 94, 96, 98, 100, 102, 104, 106, 108, 110, 112, 114, 116, 118, 120, 122, 124-131.

Also featured are pharmaceutical compositions comprising a pharmaceutically acceptable carrier and an effective amount of an anti-angiogenic peptide comprising at least about 12 amino acids of SEQ ID Nos. 2 or 4. Certain pharmaceutical compositions are comprised of an anti-angiogenic protein as disclosed herein and a second peptide. Other pharmaceutical compositions additionally comprise an effective amount of zinc. Devices, such as syringes and stents which comprise a peptide disclosed herein, are also described.

Further disclosed are nucleic acids encoding anti-angiogenic peptides, which comprise at least about 12 amino acids of SEQ ID Nos. 2 or 4, as well as pharmaceutical compositions comprising a disclosed nucleic acid in a suitable vector for expression of an effective amount of anti-angiogenic peptide to a subject. Preferred nucleic acids comprise at least about 36, 54 or 60 nucleotides of SEQ ID Nos. 1, 3 or 5. Other preferred nucleic acids are selected from the group consisting of SEQ ID Nos.: 1, 3, 5, 7, 9, 11, 13, 15, 17, 19, 21, 23, 25, 27, 29, 31, 33, 35, 37, 39, 41, 43, 45, 47, 49, 51, 53, 55, 57, 59, 61, 63, 65, 67, 69, 71, 73, 75, 77, 79, 81, 83, 85, 87, 89, 91, 93, 95, 97, 99, 101, 103, 105, 107, 109, 111, 113, 115, 117, 119, 121 and 123.

Also provided are methods for using the disclosed peptides for treating or preventing a disease or condition that results from angiogenesis (an angiogenic associated disease), such as a cancer or tumor growth.

Other features and advantages of the disclosed anti-angiogenic peptides will become apparent based on an understanding of the following detailed description and claims.

BRIEF DESCRIPTION OF THE FIGURES

FIG. 1 is a graph showing treatment of human pancreatic carcinoma (BxPC3) with human endostatin peptides.

FIGS. 2A-C are graphs showing that the N-terminal domain of endostatin is responsible for its anti-tumor properties. FIG. 2A is a graph showing treatment of LLC with murine Fc-endostatin and murine peptides P1, P2, P5, and P6 (mP1, mP2, mP5 and mP6, respectively); FIG. 2B are images of LLC sections showing CD31 staining; FIG. 2C is a graph

showing the determination of vessel density (*p<0.015 vs. PBS (control)); and FIG. 2D is a schematic diagram showing the crystal structure of endostatin.

FIG. 3A-E are graphs showing that the zinc binding site of endostatin is important for anti-tumor activity. FIG. 3A is a schematic diagram of mP1 (SEQ ID NO: 4) and mP1-H1/3A (SEQ ID NO: 150); FIG. 3B is a graph showing zinc binding to mP1 and mP1-H1/3A; Figure 3C is a graph showing treatment of LLC with mP1 and mP1-H1/3A; FIG. 3D shows images of LLC tumor sections stained with CD31; and FIG. 3E is a graph showing the determination of vessel density.

FIG. 4 is a graph showing the tumor volume of mice having LLC treated twice daily with mP1 or mP1-H with or without zinc on days 4, 7, 10 and 14.

FIG. 5 is a graph showing inhibition of endothelial cell migration by endostatin peptides.

FIGS. 6A and B show inhibition of VEGF-induced permeability by endostatin peptides. FIG. 6A is a graph showing quantification of Evan's blue dye extracted from the skin by incubation with formamide for 5 days at room temperature as measured at 620 nm; and FIG. 6B shows representative pictures of a Miles assay (V is VEGF, P is PBS).

FIG. 7 is a graph showing the tumor volume in mice to which mP1 endostatin peptides mP1, mP1-15, mP1-20 or PBS was administered as a function of days following the beginning of the treatment.

FIG. 8 shows the nucleic acid and amino acid sequence of human and murine endostatin proteins.

DETAILED DESCRIPTION

Definitions

As used herein, the following terms and phrases shall have the meanings set forth below. Unless defined otherwise, all technical and scientific terms used herein have the same meaning as commonly understood to one of ordinary skill in the art.

The singular forms "a," "an," and "the" include plural reference unless the context clearly dictates otherwise.

The term "bioavailable" when referring to a compound is art-recognized and refers to a form of a compound that allows for it, or a portion of the amount of compound administered, to be absorbed by, incorporated to, or otherwise physiologically available to a subject or patient to whom it is administered.

As used herein, the term "composition" is intended to encompass a product comprising specified ingredients in specific amounts, as well as any product which results, directly or indirectly, from combination of the specific ingredients in the specified amounts.

"Conservative substitutions" are changes between amino acids of broadly similar molecular properties. For example, interchanges within the aliphatic group alanine, valine, leucine and isoleucine can be considered as conservative. Sometimes substitution of glycine for one of these can also be considered conservative. Other conservative interchanges include those within the aliphatic group aspartate and glutamate; within the amide group asparagine and glutamine; within the hydroxyl group serine and threonine; within the aromatic group phenylalanine, tyrosine and tryptophan; within the basic group lysine, arginine and histidine; and within the sulfur-containing group methionine and cysteine. Sometimes substitution within the group methionine and leucine can also be considered conservative. Preferred conservative substitution groups are aspartate-glutamate; aspar-

agine-glutamine; valine-leucine-isoleucine; alanine-valine; phenylalanine-tyrosine; and lysine-arginine.

The terms "parenteral administration" and "administered parenterally" are art-recognized and refer to modes of administration other than enteral and topical administration, usually by injection, and includes, without limitation, intravenous, intramuscular, intraarterial, intrathecal, intracapsular, intraorbital, intracardiac, intradermal, intraperitoneal, transtracheal, subcutaneous, subcuticular, intra-articular, subcapsular, subarachnoid, intraspinal, and intrasternal injection and infusion.

A "patient," "subject" or "host" to be treated by the subject method may mean either a human or non-human animal.

The term "percent identical" refers to sequence identity between two amino acid sequences or between two nucleotide sequences. Identity can each be determined by comparing a position in each sequence which may be aligned for purposes of comparison. When an equivalent position in the compared sequences is occupied by the same base or amino acid, then the molecules are identical at that position; when the equivalent site occupied by the same or a similar amino acid residue (e.g., similar in steric and/or electronic nature), then the molecules can be referred to as homologous (similar) at that position. Expression as a percentage of homology, similarity, or identity refers to a function of the number of identical or similar amino acids at positions shared by the compared sequences. Various alignment algorithms and/or programs may be used, including FASTA, BLAST, or ENTREZ. FASTA and BLAST are available as a part of the GCG sequence analysis package (University of Wisconsin, Madison, Wis.), and can be used with, e.g., default settings. ENTREZ is available through the National Center for Biotechnology Information, National Library of Medicine, National Institutes of Health, Bethesda, Md. In one embodiment, the percent identity of two sequences can be determined by the GCG program with a gap weight of 1, e.g., each amino acid gap is weighted as if it were a single amino acid or nucleotide mismatch between the two sequences. Other techniques for alignment are described in *Methods in Enzymology*, vol. 266: Computer Methods for Macromolecular Sequence Analysis (1996), ed. Doolittle, Academic Press, Inc., a division of Harcourt Brace & Co., San Diego, Calif., USA. Preferably, an alignment program that permits gaps in the sequence is utilized to align the sequences. The Smith-Waterman is one type of algorithm that permits gaps in sequence alignments. See *Meth. Mol. Biol.* 70: 173-187 (1997). Also, the GAP program using the Needleman and Wunsch alignment method can be utilized to align sequences. An alternative search strategy uses MPSRCH software, which runs on a MASPAR computer. MPSRCH uses a Smith-Waterman algorithm to score sequences on a massively parallel computer. This approach improves ability to pick up distantly related matches, and is especially tolerant of small gaps and nucleotide sequence errors. Nucleic acid-encoded amino acid sequences can be used to search both protein and DNA databases. Databases with individual sequences are described in *Methods in Enzymology*, ed. Doolittle, supra. Databases include Genbank, EMBL, and DNA Database of Japan (DDBJ).

The term "pharmaceutically acceptable carrier" is art-recognized and refers to a pharmaceutically-acceptable material, composition or vehicle, such as a liquid or solid filler, diluent, excipient, solvent or encapsulating material, involved in carrying or transporting any subject composition or component thereof from one organ, or portion of the body, to another organ, or portion of the body. Each carrier must be "acceptable" in the sense of being compatible with the subject com-

position and its components and not injurious to the patient. Some examples of materials which may serve as pharmaceutically acceptable carriers include: (1) sugars, such as lactose, glucose and sucrose; (2) starches, such as corn starch and potato starch; (3) cellulose, and its derivatives, such as sodium carboxymethyl cellulose, ethyl cellulose and cellulose acetate; (4) powdered tragacanth; (5) malt; (6) gelatin; (7) talc; (8) excipients, such as cocoa butter and suppository waxes; (9) oils, such as peanut oil, cottonseed oil, safflower oil, sesame oil, olive oil, corn oil and soybean oil; (10) glycols, such as propylene glycol; (11) polyols, such as glycerin, sorbitol, mannitol and polyethylene glycol; (12) esters, such as ethyl oleate and ethyl laurate; (13) agar; (14) buffering agents, such as magnesium hydroxide and aluminum hydroxide; (15) alginic acid; (16) pyrogen-free water; (17) isotonic saline; (18) Ringer's solution; (19) ethyl alcohol; (20) phosphate buffer solutions; and (21) other non-toxic compatible substances employed in pharmaceutical formulations.

The terms "polynucleotide", and "nucleic acid" are used interchangeably. They refer to a polymeric form of nucleotides of any length, either deoxyribonucleotides or ribonucleotides, or analogs thereof. The following are non-limiting examples of polynucleotides: coding or non-coding regions of a gene or gene fragment, loci (locus) defined from linkage analysis, exons, introns, messenger RNA (mRNA), transfer RNA, ribosomal RNA, ribozymes, cDNA, recombinant polynucleotides, branched polynucleotides, plasmids, vectors, isolated DNA of any sequence, isolated RNA of any sequence, nucleic acid probes, and primers. A polynucleotide may comprise modified nucleotides, such as methylated nucleotides and nucleotide analogs. If present, modifications to the nucleotide structure may be imparted before or after assembly of the polymer. The sequence of nucleotides may be interrupted by non-nucleotide components. A polynucleotide may be further modified after polymerization, such as by conjugation with a labeling component. The term "recombinant" polynucleotide means a polynucleotide of genomic, cDNA, semisynthetic, or synthetic origin which either does not occur in nature or is linked to another polynucleotide in a nonnatural arrangement. The term "oligonucleotide" may be used to refer to a single stranded polynucleotide having less than about 100 nucleotides, less than about, e.g., 75, 50, 25, or 10 nucleotides.

The terms "polypeptide", "peptide" and "protein" (if single chain) are used interchangeably herein to refer to polymers of amino acids. The polymer may be linear or branched, it may comprise modified amino acids, and it may be interrupted by non-amino acids. The terms also encompass an amino acid polymer that has been modified; for example, disulfide bond formation, glycosylation, lipidation, acetylation, phosphorylation, or any other manipulation, such as conjugation with a labeling component. As used herein the term "amino acid" refers to either natural and/or unnatural or synthetic amino acids, including glycine and both the D or L optical isomers, and amino acid analogs and peptidomimetics.

The term "prophylactic" or "therapeutic" treatment is art-recognized and refers to administration of a drug to a host. If it is administered prior to clinical manifestation of the unwanted condition (e.g., disease or other unwanted state of the host animal) then the treatment is prophylactic, i.e., it protects the host against developing the unwanted condition, whereas if administered after manifestation of the unwanted condition, the treatment is therapeutic (i.e., it is intended to diminish, ameliorate or maintain the existing unwanted condition or side effects therefrom).

The term "synthetic" is art-recognized and refers to production by in vitro chemical or enzymatic synthesis.

The terms "systemic administration," "administered systemically," "peripheral administration" and "administered peripherally" are art-recognized and refer to the administration of a subject composition, therapeutic or other material other than directly into the central nervous system, such that it enters the patient's system and, thus, is subject to metabolism and other like processes.

The term "therapeutic agent" is art-recognized and refers to any chemical moiety that is a biologically, physiologically, or pharmacologically active substance that acts locally or systemically in a subject. The term thus means any substance intended for use in the diagnosis, cure, mitigation, treatment or prevention of disease or in the enhancement of desirable physical or mental development and/or conditions in an animal or human.

The term "therapeutic effect" is art-recognized and refers to a local or systemic effect in animals, particularly mammals, and more particularly humans caused by a pharmacologically active substance. The phrase "therapeutically-effective amount" means that amount of such a substance that produces some desired local or systemic effect at a reasonable benefit/risk ratio applicable to any treatment. The therapeutically effective amount of such substance will vary depending upon the subject and disease condition being treated, the weight and age of the subject, the severity of the disease condition, the manner of administration and the like, which can readily be determined by one of ordinary skill in the art. For example, certain compositions described herein may be administered in a sufficient amount to produce a at a reasonable benefit/risk ratio applicable to such treatment.

The term "treating" is art-recognized and refers to curing as well as ameliorating at least one symptom of any condition or disease or preventing a condition or disease from worsening.

The term "vector" refers to a nucleic acid capable of transporting another nucleic acid to which it has been linked. One type of vector which may be used in accord with the invention is an episome, i.e., a nucleic acid capable of extra-chromosomal replication. Other vectors include those capable of autonomous replication and expression of nucleic acids to which they are linked. Vectors capable of directing the expression of genes to which they are operatively linked are referred to herein as "expression vectors". In general, expression vectors of utility in recombinant DNA techniques are often in the form of "plasmids" which refer to circular double stranded DNA molecules which, in their vector form are not bound to the chromosome. In the present specification, "plasmid" and "vector" are used interchangeably as the plasmid is the most commonly used form of vector. However, the invention is intended to include such other forms of expression vectors which serve equivalent functions and which become known in the art subsequently hereto.

Exemplary Compositions

Peptides

Provided herein are peptides or polypeptides and variants thereof that inhibit, e.g., angiogenesis and thereby inhibit tumor growth and/or formation. Peptides may comprise an N-terminal amino acid sequence of an endostatin protein. The endostatin protein may be a mammalian protein, such as from a human, a non-human primate, a canine, a feline, an equine, a bovine, an ovine, a sheep, or a rodent (e.g., mouse or rat). A nucleotide sequence encoding human endostatin is set forth in SEQ ID NO: 151 and the protein encoded thereby is set forth as SEQ ID NO: 152 and is identical to GenBank Accession number AAF01310 except that it is lacking the initiator

methionine of AAF01310. A nucleotide sequence encoding mouse endostatin is set forth in SEQ ID NO: 153 and the protein encoded thereby is set forth as SEQ ID NO: 154 and is identical to GenBank Accession number AAF69009. The nucleotide and amino acid sequences of other species are also publicly available.

In one embodiment, the peptide comprises about 5 to about 40 amino acids of the N-terminal region of an endostatin protein. The peptide may comprise from about 10 to about 30 or 35 amino acids or from about 20 to about 30 or 35 amino acids of the N-terminal region of an endostatin protein. For example, a peptide may comprise about amino acids 1 to about amino acids 35, 30, 25, 20, 15, or 10 of an endostatin protein, such as a protein having SEQ ID NO: 2 or 4. The amino acid sequences of these human and mouse peptides are HSHRDFQPVHLHLVALNSPLSGGMRGIR (SEQ ID NO: 2) and HTHQDFQPVHLHLVALNTPLSGGMRGIR (mP1; SEQ ID NO: 4), respectively. SEQ ID NO: 2 is encoded by the following nucleic acid sequence: cacagccaccgagcttcagccg-gtgcctccacctgggtgcgctcaacagc-cccctgtcaggcggcatgcggggcatccgc (SEQ ID NO: 1). SEQ ID NO: 4 is encoded by the following nucleic acid sequence: catactcatcaggactttcagccagt-gctccacctgggtgcactgaacacccctgtctggagggcatgcgtggtatccgt (SEQ ID NO: 3).

Peptides, which are slightly different from SEQ ID NO: 1 have been shown to retain anti-angiogenic activity. One such peptide is HSHRDFQPVHLHLVALNSPLSGGMRG (hP1; SEQ ID NO: 6), which is encoded by the nucleic acid sequence: catactcatcaggactttcagccagt-gctccacctgggtgcactgaacacccctgtctggagggcatgcgtggt (SEQ ID NO: 5). SEQ ID NO: 6 does not contain the two most C-terminal amino acid residues in SEQ ID NO: 2. Further anti-angiogenic peptides may lack one or more amino acids at the N— or C-terminus of SEQ ID Nos. 2 or 4. Exemplary anti-angiogenic peptides are as follows:

SHRDFQPVHLHLVALNSPLSGGMRGIR;	(SEQ ID NO: 8)
HRDFQPVHLHLVALNSPLSGGMRGIR;	(SEQ ID NO: 10)
RDFQPVHLHLVALNSPLSGGMRGIR;	(SEQ ID NO: 12)
DFQPVHLHLVALNSPLSGGMRGIR;	(SEQ ID NO: 14)
FQPVHLHLVALNSPLSGGMRGIR;	(SEQ ID NO: 16)
QPVHLHLVALNSPLSGGMRGIR;	(SEQ ID NO: 18)
PVHLHLVALNSPLSGGMRGIR;	(SEQ ID NO: 20)
VLHLVALNSPLSGGMRGIR;	(SEQ ID NO: 22)
LHLVALNSPLSGGMRGIR;	(SEQ ID NO: 24)
HLVALNSPLSGGMRGIR;	(SEQ ID NO: 26)
LVALNSPLSGGMRGIR;	(SEQ ID NO: 28)
VALNSPLSGGMRGIR;	(SEQ ID NO: 30)
ALNSPLSGGMRGIR;	(SEQ ID NO: 32)
LNSPLSGGMRGIR;	(SEQ ID NO: 34)
NSPLSGGMRGIR;	(SEQ ID NO: 36)
HSHRDFQPVHLHLVALNSPLSGGMRGI;	(SEQ ID NO: 38)
HSHRDFQPVHLHLVALNSPLSGGMR;	(SEQ ID NO: 40)
HSHRDFQPVHLHLVALNSPLSGGM;	(SEQ ID NO: 42)

-continued

HSHRDFQPVHLHLVALNSPLSGG;	(SEQ ID NO: 44)
5 HSHRDFQPVHLHLVALNSPLSG;	(SEQ ID NO: 46)
HSHRDFQPVHLHLVALNSPLS;	(SEQ ID NO: 48)
HSHRDFQPVHLHLVALNSPL;	(SEQ ID NO: 50)
10 HSHRDFQPVHLHLVALNSP;	(SEQ ID NO: 52)
HSHRDFQPVHLHLVALNS;	(SEQ ID NO: 54)
HSHRDFQPVHLHLVALN;	(SEQ ID NO: 56)
15 HSHRDFQPVHLHLVAL;	(SEQ ID NO: 58)
HSHRDFQPVHLHLVA;	(SEQ ID NO: 60)
HSHRDFQPVHLHLV;	(SEQ ID NO: 62)
20 HSHRDFQPVHLHL;	(SEQ ID NO: 64)
THQDFQPVHLHLVALNTPLSGGMRGIR;	(SEQ ID NO: 66)
HQDFQPVHLHLVALNTPLSGGMRGIR;	(SEQ ID NO: 68)
25 QDFQPVHLHLVALNTPLSGGMRGIR;	(SEQ ID NO: 70)
DFQPVHLHLVALNTPLSGGMRGIR;	(SEQ ID NO: 72)
FQPVHLHLVALNTPLSGGMRGIR;	(SEQ ID NO: 74)
QPVHLHLVALNTPLSGGMRGIR;	(SEQ ID NO: 76)
30 PVHLHLVALNTPLSGGMRGIR;	(SEQ ID NO: 78)
VLHLVALNTPLSGGMRGIR;	(SEQ ID NO: 80)
LHLVALNTPLSGGMRGIR;	(SEQ ID NO: 82)
35 HLVALNTPLSGGMRGIR;	(SEQ ID NO: 84)
LVALNTPLSGGMRGIR;	(SEQ ID NO: 86)
VALNTPLSGGMRGIR;	(SEQ ID NO: 88)
40 ALNTPLSGGMRGIR;	(SEQ ID NO: 90)
LNTPLSGGMRGIR;	(SEQ ID NO: 92)
NTPLSGGMRGIR;	(SEQ ID NO: 94)
45 HTHQDFQPVHLHLVALNTPLSGGMRGI;	(SEQ ID NO: 96)
HTHQDFQPVHLHLVALNTPLSGGMRG;	(SEQ ID NO: 98)
HTHQDFQPVHLHLVALNTPLSGGMR;	(SEQ ID NO: 100)
50 HTHQDFQPVHLHLVALNTPLSGGM;	(SEQ ID NO: 102)
HTHQDFQPVHLHLVALNTPLSGG;	(SEQ ID NO: 104)
HTHQDFQPVHLHLVALNTPLSG;	(SEQ ID NO: 106)
55 HTHQDFQPVHLHLVALNTPLS;	(mP1-20; SEQ ID NO: 108)
HTHQDFQPVHLHLVALNTPL;	(SEQ ID NO: 110)
HTHQDFQPVHLHLVALNTPL;	(SEQ ID NO: 112)
60 HTHQDFQPVHLHLVALNT;	(SEQ ID NO: 114)
HTHQDFQPVHLHLVALN;	(SEQ ID NO: 116)
HTHQDFQPVHLHLVAL;	(mP1-15; SEQ ID NO: 118)
HTHQDFQPVHLHLVA;	(SEQ ID NO: 120)
65 HTHQDFQPVHLHLV;	(SEQ ID NO: 122)

-continued

HTHQDFQPVHLHL; (SEQ ID NO: 124)
 HSHRDFVALNSPLSGGMRGIR; (SEQ ID NO: 125)
 HSHRDFQPVHLHLLSGGMRGIR; (SEQ ID NO: 126)
 QPVLHLVALNTPLSGGMRGIR; (SEQ ID NO: 127)
 HTHQDFVALNTPLSGGMRGIR; (SEQ ID NO: 128)
 and
 HTHQDFQPVHLHLLSGGMRGIR. (SEQ ID NO: 129)

Other anti-angiogenic peptides are based on the following consensus sequences: HXaaHXaaDFQPVHLVALNXaa-PLSGGMRGIR (SEQ ID NO: 130) or HXaaHXaaDFQPV-LHLVALNXaaPLSG (SEQ ID NO: 131), wherein Xaa is any amino acid.

Other peptides having anti-angiogenic activity may comprise, consist of or consist essentially of any of the amino acid sequences described above. Yet other peptides may comprise, consist of or consist essentially of an amino acid sequence that has at least about 70%, 80%, 90%, 95%, 98% or 99% identity or homology with an N-terminal endostatin peptide. For example, peptides that differ from a sequence in a naturally occurring endostatin protein in about 1, 2, 3, 4, 5 or more amino acids would be expected to retain anti-angiogenic activity. Peptides that are similar to the sequences described above may contain substitutions, e.g., conservative substitutions, deletions or additions. The differences are preferably in regions that are not significantly conserved among different species. Such regions can be identified by aligning the amino acid sequences of endostatin proteins from various animal species. For example, amino acids 2, 4, and 17 of SEQ ID NO: 2, 4, or 6, and the highlighted amino acids in e.g. SEQ ID NOs: 8, 10, 12, 30, 32, or 34 may be substituted without negatively impacting the anti-angiogenic activity, since these amino acids differ in the human and mouse sequences. These amino acids can be substituted, e.g., with those found in another species. Amino acid 9 can also be substituted, since that one is different in the *Gallus gallus* species. Other amino acids that may be substituted, inserted or deleted at these or other locations can be identified by mutagenesis studies coupled with biological assays.

Also encompassed herein are endostatin peptides that are fused to a heterologous peptide, such as a peptide that can be used for detecting; purifying; stabilizing; or solubilizing the endostatin peptide.

A peptide may be linked to an immunoglobulin (Ig) constant heavy or light chain domain or portion thereof. For example, a peptide may be linked to a CH1, CH2 and/or CH3 domain of a heavy chain. If the constant region is from a light chain, it may be from a kappa or lambda light chain. If the constant region is from a heavy chain, it may be from an antibody of any one of the following classes of antibodies: IgG, IgA, IgE, IgD, and IgM. IgG may be IgG1, IgG2, IgG3 or IgG4. The constant domain may be an Fc fragment. The constant domain may be from a mammalian antibody, e.g., a human antibody. Soluble receptor-IgG fusion proteins are common immunological reagents and methods for their construction are known in the art (see e.g., U.S. Pat. Nos. 5,225, 538, 5,726,044; 5,707,632; 750,375, 5,925,351, 6,406,697 and Bergers et al. *Science* 1999 284: 808-12). Preferred as immunoglobulin is the constant part of the heavy chain of human IgG, particularly IgG1, where dimerization between two heavy chains takes place at the hinge region. It is recognized that inclusion of the CH2 and CH3 domains of the Fc region as part of the fusion polypeptide increases the in vivo

circulation half-life of the polypeptide comprising the Fc region, and that of the oligomer or dimer comprising the polypeptide.

An Fc portion of human IgG1 which includes the hinge region, and domains CH2 and CH3 has the nucleotide sequence 5' gag ccc aaa tct tgt gac aaa act cac aca tgc cca ccg tgc cca gca cct gaa ctc ctg ggg gga ccg tca gtc ttc ctc ttc ccc cca aaa ccc aag gac acc ctc atg atc tcc cgg acc cct gag gtc aca tgc gtg gtg gtg gac gtg agc cac gaa gac cct gag gtc aag ttc aac tgg tac gtg gac ggc gtg gag gtg cat aat gcc aag aca aag ccg cgg gag gag cag tac aac agc acg tac cgt gtg gtc agc gtc ctc acc gtc ctg cac cag gac tgg ctg aat ggc aag gag tac aag tgc aag gtc tcc aac aaa gcc ctc cca gcc ccc atc gag aaa acc atc tcc aaa gcc aaa ggg cag ccc cga gaa cca cag gtg tac acc ctg ccc cca tcc cgg gat gag ctg acc aag aac cag gtc agc ctg acc tgc ctg gtc aaa ggc ttc tat ccc agc gac atc gcc gtg gag tgg gag agc aat ggg cag ccg gag aac aac tac aag acc acg cct ccc gtg ctg gac tcc gac ggc tcc ttc ttc ctc tac agc aag ctc acc gtg gac aag agc agg tgg cag cag ggg aac gtc ttc tca tgc tcc gtg atg cat gag gct ctg cac aac cac tac acg cag aag agc ctc tcc ctg tct ccg ggt aaa tga 3' (SEQ ID NO: 132), which encodes a peptide having the amino acid sequence: Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gin Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gin Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gin Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gin Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gin Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gin Lys Ser Leu Ser Leu Ser Pro Gly Lys (SEQ ID NO: 133).

Constant Ig domains may also contain one or more mutations that reduce or eliminate one or more effector function, e.g., binding to Fc receptors and complement activation (see, e.g., S. Morrison, *Annu. Rev. Immunol.*, 10, pp. 239-65 (1992); Duncan and Winter (1988) *Nature* 332: 738-740; and Xu et al. (1994) *J Biol. Chem.* 269: 3469-3474). For example, mutations of amino acids corresponding to Leu 235 and Pro 331 of human IgG1 to Glu and Ser respectively, are provided. Such constructs are further described in U.S. Pat. No. 6,656, 728.

The constant Ig domain may be linked to the N-terminus or C-terminus of a peptide.

The peptide may also be linked to a linker sequence with a thrombin cleavage site, such as between the peptide and an immunoglobulin domain. An exemplary nucleotide sequence encoding such a site has the following nucleotide sequence: 5' tct aga ggt ggt cta gtg ccg cgc ggc agc ggt tcc ccc ggg ttg cag 3' (SEQ ID NO: 134), which encodes a peptide having the amino acid sequence: Ser Arg Gly Gly Leu Val Pro Arg Gly Ser Gly Ser Pro Gly Leu Gin (SEQ ID NO: 135).

A peptide may also be fused to a signal sequence. For example, when prepared recombinantly, a nucleic acid encoding the peptide may be linked at its 5' end to a signal sequence, such that the peptide is secreted from the cell.

Peptides may be used as a substantially pure preparation, e.g., wherein at least about 90% of the peptides in the preparation are the desired peptide. Compositions comprising at least about 50%, 60%, 70%, or 80% of the desired peptide may also be used.

Peptides may be denatured or non-denatured and may be aggregated or non-aggregated as a result thereof. Peptides can be denatured according to methods known in the art.

Peptides may be conjugated to zinc. Thus, peptides may be in a composition comprising Zn^{2+} , e.g., in sufficient quanti-

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ties that most of the peptides are conjugated to one or more Zn^{2+} molecule. Binding of Zn^{2+} to a peptide can be demonstrated by the following assay. Zinc and peptide solutions are mixed, optionally incubated together, and then dialyzed to remove the zinc that is not bound to the peptides. Detection of zinc in the peptide solution can then be performed by atomic absorption.

Yet other peptides that are encompassed herein are those that comprise modified amino acids. Exemplary peptides are derivative peptides that may be one modified by glycosylation, pegylation, phosphorylation or any similar process that retains at least one biological function of the peptide from which it was derived.

Peptides may also comprise one or more non-naturally occurring amino acids. For example, nonclassical amino acids or chemical amino acid analogs can be introduced as a substitution or addition into peptides. Non-classical amino acids include, but are not limited to, the D-isomers of the common amino acids, 2,4-diaminobutyric acid, alpha-amino isobutyric acid, 4-aminobutyric acid, Abu, 2-amino butyric acid, gamma-Abu, epsilon-Ahx, 6-amino hexanoic acid, Aib, 2-amino isobutyric acid, 3-amino propionic acid, ornithine, norleucine, norvaline, hydroxyproline, sarcosine, citrulline, homocitrulline, cysteic acid, t-butylglycine, t-butylalanine, phenylglycine, cyclohexylalanine, beta-alanine, fluoro-amino acids, designer amino acids such as beta-methyl amino acids, Calpha-methyl amino acids, Nalpha-methyl amino acids, and amino acid analogs in general. Furthermore, the amino acid can be D (dextrorotary) or L (levorotary).

In other specific embodiments, branched versions of the peptides listed herein are provided, e.g., by substituting one or more amino acids within the sequence with an amino acid or amino acid analog with a free side chain capable of forming a peptide bond with one or more amino acids (and thus capable of forming a "branch"). Cyclical peptides are also contemplated.

Also included are peptide derivatives which are differentially modified during or after synthesis, e.g., by benzylation, glycosylation, acetylation, phosphorylation, amidation, pegylation, derivatization by known protecting/blocking groups, proteolytic cleavage, linkage to an antibody molecule or other cellular ligand, etc. In specific embodiments, the peptides are acetylated at the N-terminus and/or amidated at the C-terminus.

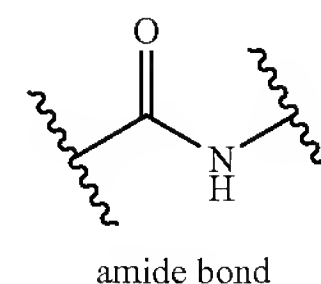
Also provided are derivatives of endostatin peptides, such as chemically modified peptides and peptidomimetics. Peptidomimetics are compounds based on, or derived from, peptides and proteins. Peptidomimetics can be obtained by structural modification of known peptide sequences using unnatural amino acids, conformational restraints, isosteric replacement, and the like. The subject peptidomimetics constitute the continuum of structural space between peptides and non-peptide synthetic structures; peptidomimetics may be useful, therefore, in delineating pharmacophores and in helping to translate peptides into nonpeptide compounds with the activity of the parent peptides.

Moreover, mimetopes of the subject peptides can be provided. Such peptidomimetics can have such attributes as being non-hydrolyzable (e.g., increased stability against proteases or other physiological conditions which degrade the corresponding peptide), increased specificity and/or potency for stimulating cell differentiation. For illustrative purposes, peptide analogs can be generated using, for example, benzodiazepines (e.g., see Freidinger et al. in *Peptides: Chemistry and Biology*, G. R. Marshall ed., ESCOM Publisher: Leiden, Netherlands, 1988), substituted gamma lactam rings (Garvey et al. in *Peptides: Chemistry and Biology*, G. R. Marshall ed., ESCOM Publisher: Leiden, Netherlands, 1988, p123), C-7 mimics (Huffman et al. in *Peptides: Chemistry and Biology*, G. R. Marshall ed., ESCOM Publisher: Leiden, Netherlands, 1988, p. 105), keto-methylene pseudopeptides (Ewenson et

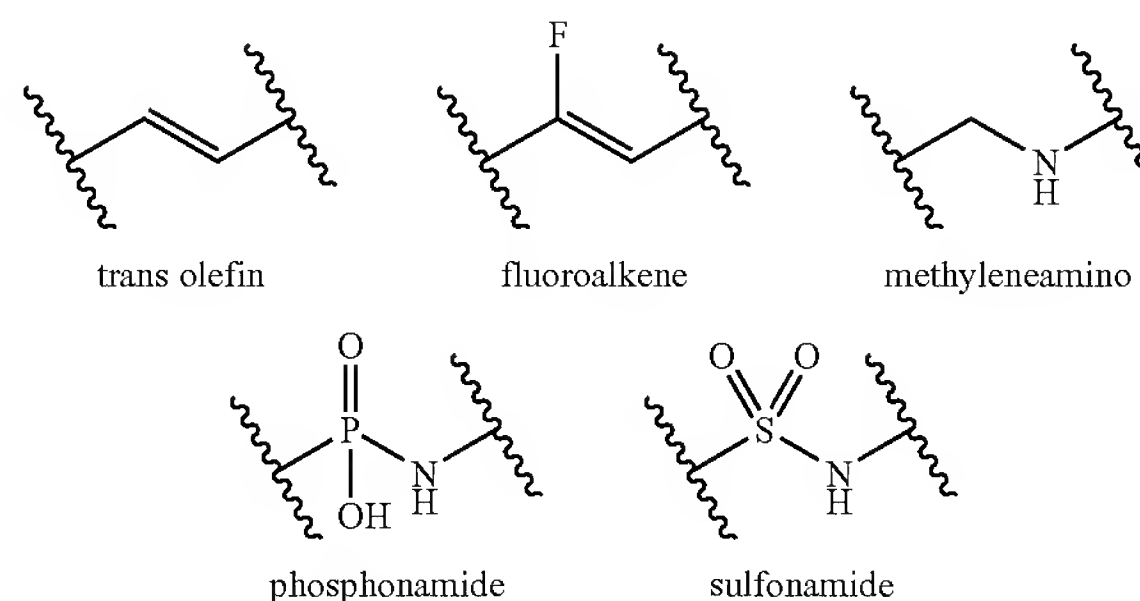
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al. (1986) *J Med Chem* 29:295; and Ewenson et al. in *Peptides: Structure and Function* (Proceedings of the 9th American Peptide Symposium) Pierce Chemical Co. Rockland, Ill., 1985), β -turn dipeptide cores (Nagai et al. (1985) *Tetrahedron Lett* 26:647; and Sato et al. (1986) *J Chem Soc Perkin Trans 1*: 1231), β -aminoalcohols (Gordon et al. (1985) *Biochem Biophys Res Commun* 126:419; and Dann et al. (1986) *Biochem Biophys Res Commun* 134:71), diaminoketones (Natarajan et al. (1984) *Biochem Biophys Res Commun* 124: 141), and methyleneamino-modified (Roark et al. in *Peptides: Chemistry and Biology*, G. R. Marshall ed., ESCOM Publisher: Leiden, Netherlands, 1988, p134). Also, see generally, Session III: Analytic and synthetic methods, in in *Peptides: Chemistry and Biology*, G. R. Marshall ed., ESCOM Publisher: Leiden, Netherlands, 1988).

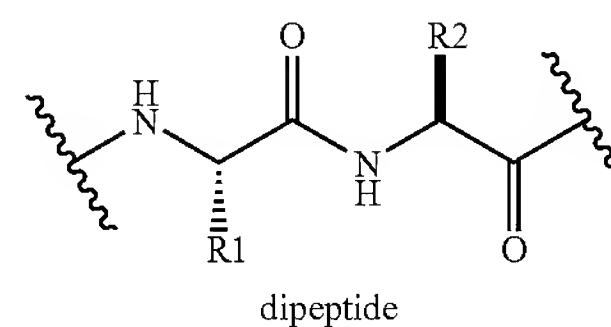
In addition to a variety of sidechain replacements which can be carried out to generate peptidomimetics, the description specifically contemplates the use of conformationally restrained mimics of peptide secondary structure. Numerous surrogates have been developed for the amide bond of peptides. Frequently exploited surrogates for the amide bond include the following groups (i) trans-olefins, (ii) fluoroalkene, (iii) methyleneamino, (iv) phosphonamides, and (v) sulfonamides.



Examples of Surrogates:

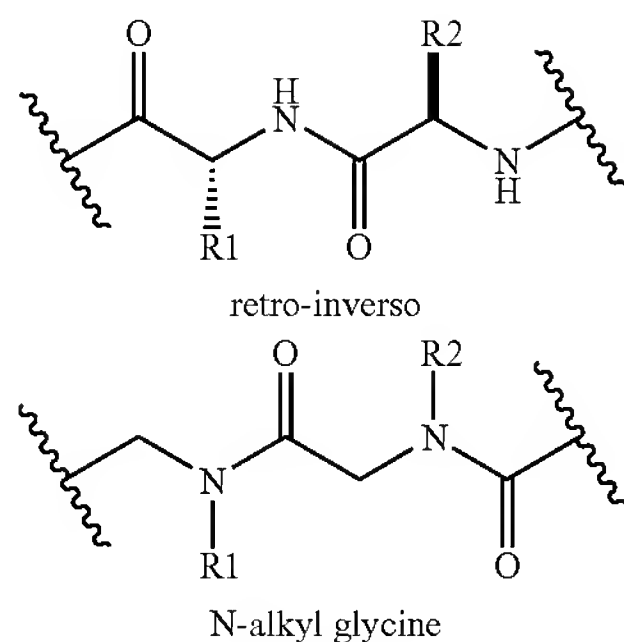


Additionally, peptidomimetics based on more substantial modifications of the backbone of a peptide can be used. Peptidomimetics which fall in this category include (i) retro-inverso analogs, and (ii) N-alkyl glycine analogs (so-called peptoids).

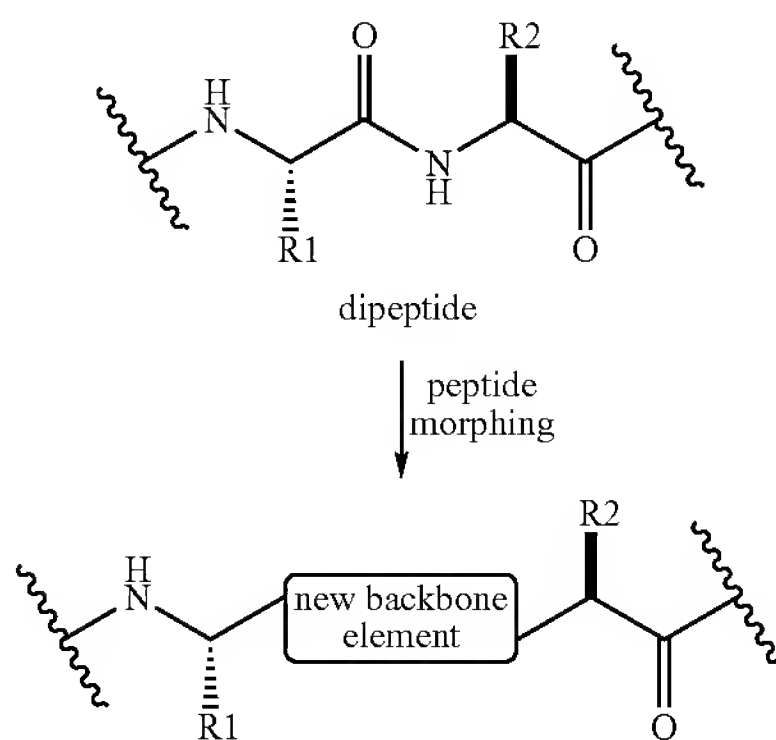


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Examples of analogs:



Furthermore, the methods of combinatorial chemistry are being brought to bear, on the development of new peptidomimetics. For example, one embodiment of a so-called "peptide morphing" strategy focuses on the random generation of a library of peptide analogs that comprise a wide range of peptide bond substitutes.



In an exemplary embodiment, the peptidomimetic can be derived as a retro-inverso analog of the peptide. Such retro-inverso analogs can be made according to the methods known in the art, such as that described by the Sisto et al. U.S. Pat. No. 4,522,752. A retro-inverso analog can be generated as described, e.g., in WO 00/01720. It will be understood that a mixed peptide, e.g. including some normal peptide linkages, may be generated. As a general guide, sites which are most susceptible to proteolysis are typically altered, with less susceptible amide linkages being optional for mimetic switching. The final product, or intermediates thereof, can be purified by HPLC.

Peptides may comprise at least one amino acid or every amino acid that is a D stereoisomer. Other peptides may comprise at least one amino acid that is reversed. The amino acid that is reversed may be a D stereoisomer. Every amino acid of a peptide may be reversed and/or every amino acid may be a D stereoisomer.

In another illustrative embodiment, a peptidomimetic can be derived as a retro-enantio analog of a peptide. Retro-enantio analogs such as this can be synthesized with commercially available D-amino acids (or analogs thereof) and standard solid- or solution-phase peptide-synthesis techniques, as described, e.g., in WO 00/01720. The final product may be purified by HPLC to yield the pure retro-enantio analog.

In still another illustrative embodiment, trans-olefin derivatives can be made for the subject peptide. Trans-olefin analogs can be synthesized according to the method of Y. K. Shue et al. (1987) *Tetrahedron Letters* 28:3225 and as

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described in WO 00/01720. It is further possible to couple pseudopeptides synthesized by the above method to other pseudopeptides, to make peptide analogs with several olefinic functionalities in place of amide functionalities.

Still another class of peptidomimetic derivatives include the phosphonate derivatives. The synthesis of such phosphonate derivatives can be adapted from known synthesis schemes. See, for example, Loots et al. in *Peptides: Chemistry and Biology*, (Escom Science Publishers, Leiden, 1988, p. 118); Petrillo et al. in *Peptides: Structure and Function* (Proceedings of the 9th American Peptide Symposium, Pierce Chemical Co. Rockland, Ill., 1985).

Many other peptidomimetic structures are known in the art and can be readily adapted for use in the subject peptidomimetics. To illustrate, a peptidomimetic may incorporate the 1-azabicyclo[4.3.0]nonane surrogate (see Kim et al. (1997) *J. Org. Chem.* 62:2847), or an N-acyl piperazic acid (see Xi et al. (1998) *J. Am. Chem. Soc.* 120:80), or a 2-substituted piperazine moiety as a constrained amino acid analogue (see Williams et al. (1996) *J. Med. Chem.* 39:1345-1348). In still other embodiments, certain amino acid residues can be replaced with aryl and bi-aryl moieties, e.g., monocyclic or bicyclic aromatic or heteroaromatic nucleus, or a biaromatic, aromatic-heteroaromatic, or biheteroaromatic nucleus.

The subject peptidomimetics can be optimized by, e.g., combinatorial synthesis techniques combined with high throughput screening.

Moreover, other examples of mimetopes include, but are not limited to, protein-based compounds, carbohydrate-based compounds, lipid-based compounds, nucleic acid-based compounds, natural organic compounds, synthetically derived organic compounds, anti-idiotypic antibodies and/or catalytic antibodies, or fragments thereof. A mimetope can be obtained by, for example, screening libraries of natural and synthetic compounds for compounds capable of inhibiting angiogenesis and/or tumor growth. A mimetope can also be obtained, for example, from libraries of natural and synthetic compounds, in particular, chemical or combinatorial libraries (i.e., libraries of compounds that differ in sequence or size but that have the same building blocks). A mimetope can also be obtained by, for example, rational drug design. In a rational drug design procedure, the three-dimensional structure of a compound of the present invention can be analyzed by, for example, nuclear magnetic resonance (NMR) or x-ray crystallography. The three-dimensional structure can then be used to predict structures of potential mimetopes by, for example, computer modelling. The predicted mimetope structures can then be produced by, for example, chemical synthesis, recombinant DNA technology, or by isolating a mimetope from a natural source (e.g., plants, animals, bacteria and fungi).

"Peptides, variants and derivatives thereof" or "peptides and analogs thereof" are included in "peptide therapeutics" and is intended to include any of the peptides or modified forms thereof, e.g., peptidomimetics, described herein. Preferred peptide therapeutics have anti-angiogenic activity. For example, they may reduce or inhibit angiogenesis by a factor of at least about 50%, 2 fold, 5 fold, 10 fold, 30 fold or 100 fold, as determined, e.g., in an assay described herein.

Assays for testing candidate peptides for anti-angiogenesis and inhibition of tumor growth or formation are known in the art and exemplary ones are further described herein.

Nucleic Acids

Also disclosed are nucleic acids encoding anti-angiogenic peptides. Preferred nucleic acids are as follows:

cacagccaccgcgacttccagccggtgctccacctggttgcgctcaacagccccctgtcaggcggcacatgcggggcat
 ccgc. (SEQ ID NO: 1)

catactcatcaggactttcagccagtgtccacctggtggcactgaacacccccctgtctggaggcatgcgtggtatccg
 t. (SEQ ID NO: 3)

catactcatcaggactttcagccagtgtccacctggtggcactgaacacccccctgtctggaggcatgcgtggt. (SEQ ID NO: 5)

agccaccgcgacttccagccggtgctccacctggttgcgctcaacagccccctgtcaggcggcacatgcggggcatcc
 gc; (SEQ ID NO: 7)

caccgcgacttccagccggtgctccacctggttgcgctcaacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 9)

cgcgacttccagccggtgctccacctggttgcgctcaacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 11)

gacttccagccggtgctccacctggttgcgctcaacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 13)

ttccagccggtgctccacctggttgcgctcaacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 15)

cagccggtgctccacctggttgcgctcaacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 17)

ccggtgctccacctggttgcgctcaacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 19)

gtgctccacctggttgcgctcaacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 21)

ctccacctggttgcgctcaacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 23)

cacctggttgcgctcaacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 25)

ctggttgcgctcaacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 27)

gttgctcaacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 29)

gcgctcaacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 31)

ctcaacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 33)

aacagccccctgtcaggcggcacatgcggggcatccgc; (SEQ ID NO: 35)

cacagccaccgcgacttccagccggtgctccacctggttgcgctcaacagccccctgtcaggcggcacatgcggggcat
 c; (SEQ ID NO: 37)

cacagccaccgcgacttccagccggtgctccacctggttgcgctcaacagccccctgtcaggcggcacatgcggg; (SEQ ID NO: 39)

cacagccaccgcgacttccagccggtgctccacctggttgcgctcaacagccccctgtcaggcggcacatg; (SEQ ID NO: 41)

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cacagccaccgcgacttccagccggtgctccacctggttgcgctcaacagccccctgtca; (SEQ ID NO: 47)

cacagccaccgcgacttccagccggtgctccacctggttgcgctcaacagccccctg; (SEQ ID NO: 49)

cacagccaccgcgacttccagccggtgctccacctggttgcgctcaacagcccc; (SEQ ID NO: 51)

cacagccaccgcgacttccagccggtgctccacctggttgcgctcaacagc; (SEQ ID NO: 53)

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cagccagtgtccacctggtggcactgaacacccccctgtctggaggcatgcgtggtatccgt; (SEQ ID NO: 75)

- continued

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gtgtccacctggtggcactgaacacccccctgtctggaggcatgcgtggtatccgt; (SEQ ID NO: 79)

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catactcatcaggactttcagccagtgtccacctggtggcactgaac; (SEQ ID NO: 115)

catactcatcaggactttcagccagtgtccacctggtggcactg; (SEQ ID NO: 117)

catactcatcaggactttcagccagtgtccacctggtggca; (SEQ ID NO: 119)

catactcatcaggactttcagccagtgtccacctggtg; (SEQ ID NO: 121)

and

catactcatcaggactttcagccagtgtccacctg. (SEQ ID NO: 123)

Nucleic acids include vectors, such as expression vectors for producing a peptide, e.g., viral vectors. Also encompassed herein are cells comprising a nucleic acid encoding a peptide described herein and methods for producing peptides comprising culturing these cells to produce a peptide. These methods can be used of producing recombinant peptides or for expression of a peptide in a cell, e.g., in a cell of a subject.

Appropriate vectors may be introduced into host cells using well known techniques such as infection, transduction, transfection, transvection, electroporation and transformation. The vector may be, for example, a phage, plasmid, viral or retroviral vector. Retroviral vectors may be replication competent or replication defective. In the latter case, viral propagation generally will occur only in complementing host cells.

The vector may contain a selectable marker for propagation in a host. Generally, a plasmid vector is introduced in a precipitate, such as a calcium phosphate precipitate, or in a complex with a charged lipid. If the vector is a virus, it may be packaged in vitro using an appropriate packaging cell line and then transduced into host cells.

Preferred vectors comprise cis-acting control regions to the polynucleotide of interest. Appropriate trans-acting factors

may be supplied by the host, supplied by a complementing vector or supplied by the vector itself upon introduction into the host.

In certain embodiments, the vectors provide for specific expression, which may be inducible and/or cell type-specific. Particularly preferred among such vectors are those inducible by environmental factors that are easy to manipulate, such as temperature and nutrient additives.

Expression vectors useful in the present invention include chromosomal-, episomal- and virus-derived vectors, e.g., vectors derived from bacterial plasmids, bacteriophage, yeast episomes, yeast chromosomal elements, viruses such as baculoviruses, papova viruses, vaccinia viruses, adenoviruses, fowl pox viruses, pseudorabies viruses and retroviruses, and vectors derived from combinations thereof, such as cosmids and phagemids.

The DNA insert should be operatively linked to an appropriate promoter, such as the phage lambda PL promoter, the *E. coli* lac, trp and tac promoters, the SV40 early and late promoters and promoters of retroviral LTRs, to name a few. Other suitable promoters will be known to the skilled artisan. The expression constructs will further contain sites for transcription initiation, termination and, in the transcribed region, a ribosome binding site for translation. The coding portion of

the mature transcripts expressed by the constructs will preferably include a translation initiating site at the beginning and a termination codon (UAA, UGA or UAG) appropriately positioned at the end of the polypeptide to be translated.

As indicated, the expression vectors will preferably include at least one selectable marker. Such markers include dihydrofolate reductase or neomycin resistance for eukaryotic cell culture and tetracycline, kanamycin, or ampicillin resistance genes for culturing in *E. coli* and other bacteria. Representative examples of appropriate hosts include, but are not limited to, bacterial cells, such as *E. coli*, *Streptomyces* and *Salmonella typhimurium* cells; fungal cells, such as yeast cells; insect cells such as *Drosophila* S2 and Sf9 cells; animal cells such as CHO, COS and Bowes melanoma cells; and plant cells. Appropriate culture mediums and conditions for the above-described host cells are known in the art.

Among vectors preferred for use in bacteria include pQE70, pQE60 and pQE9, pQE10 available from Qiagen; pBS vectors, Phagescript vectors, Bluescript vectors, pNH8A, pNH16a, pNH18A, pNH46A available from Stratagene; pET series of vectors available from Novagen; and ptrc99a, pKK223-3, pKK233-3, pDR540, pRIT5 available from Pharmacia. Among preferred eukaryotic vectors are pWLNEO, pSV2CAT, pOG44, pXT1 and pSG available from Stratagene; and pSVK3, pBPV, pMSG and pSVL available from Pharmacia. Other suitable vectors will be readily apparent to the skilled artisan.

Among known bacterial promoters suitable for use in the present invention include the *E. coli* lacI and lacZ promoters, the T3, T5 and T7 promoters, the gpt promoter, the lambda PR and PL promoters, the trp promoter and the xyl/tet chimeric promoter. Suitable eukaryotic promoters include the CMV immediate early promoter, the HSV thymidine kinase promoter, the early and late SV40 promoters, the promoters of retroviral LTRs, such as those of the Rous sarcoma virus (RSV), and metallothionein promoters, such as the mouse metallothionein-I promoter.

Introduction of the construct into the host cell can be effected by calcium phosphate transfection, DEAE-dextran mediated transfection, cationic lipid-mediated transfection, electroporation, transduction, infection or other methods. Such methods are described in many standard laboratory manuals (for example, Davis, et al., *Basic Methods In Molecular Biology* (1986)).

Transcription of DNA encoding the polypeptides of the present invention by higher eukaryotes may be increased by inserting an enhancer sequence into the vector. Enhancers are cis-acting elements of DNA, usually about from 10 to 300 nucleotides that act to increase transcriptional activity of a promoter in a given host cell-type. Examples of enhancers include the SV40 enhancer, which is located on the late side of the replication origin at nucleotides 100 to 270, the cytomegalovirus early promoter enhancer, the polyoma enhancer on the late side of the replication origin, and adenovirus enhancers.

A recombinant soluble form of a polypeptide of the present invention may also be produced, e.g., by deleting at least a portion of the transmembrane domain, such that the protein is not capable to localize itself to a cell membrane. Also within the scope of the invention are nucleic acids encoding splice variants or nucleic acids representing transcripts synthesized from an alternative transcriptional initiation site, such as those whose transcription was initiated from a site in an intron. Such homologues can be cloned by hybridization or PCR using standard methods known in the art.

The polynucleotide sequence may also encode for a leader sequence, e.g., the natural leader sequence or a heterologous

leader sequence. Alternatively, the nucleic acid can be engineered such that the natural leader sequence is deleted and a heterologous leader sequence inserted in its place. The term "leader sequence" is used interchangeably herein with the term "signal peptide". For example, the desired DNA sequence may be fused in the same reading frame to a DNA sequence which aids in expression and secretion of the polypeptide from the host cell, for example, a leader sequence which functions as a secretory sequence for controlling transport of the polypeptide from the cell. The protein having a leader sequence is a preprotein and may have the leader sequence cleaved by the host cell to form the mature form of the protein.

For secretion of the translated polypeptide into the lumen of the endoplasmic reticulum, into the periplasmic space or into the extracellular environment, appropriate secretion signals may be incorporated into the expressed polypeptide, for example, the amino acid sequence KDEL. The signals may be endogenous to the polypeptide or they may be heterologous signals.

The polypeptides may be expressed in a modified form, such as a fusion protein, and may include not only secretion signals, but also additional heterologous functional regions. For instance, a region of additional amino acids, particularly charged amino acids, may be added to the N-terminus or C-terminus of the polypeptide to improve stability and persistence in the host cell, during purification, or during subsequent handling and storage. Also, peptide moieties may be added to the polypeptide to facilitate purification. Such regions may be removed prior to final preparation of the polypeptide. The addition of peptide moieties to polypeptides to engender secretion or excretion, to improve stability and to facilitate purification, among others, are familiar and routine techniques in the art. An example of such a fusion protein may comprise a heterologous region from immunoglobulin that is useful to solubilize proteins.

Exemplary Methods

Anti-angiogenic peptides or an anti-angiogenic peptide encoding nucleic acid vector may be administered to a subject in need thereof to prevent or reduce angiogenesis and/or cell proliferation, e.g., tumor growth, or any disease or disorder associated therewith. The subject may be a human or animal, such as a mammal.

Unregulated angiogenesis occurs in a multiplicity of disease states, tumor metastasis and abnormal growth by endothelial cells and supports the pathological damage seen in these conditions. The diverse pathological states created due to unregulated angiogenesis have been grouped together as angiogenic dependent or angiogenic associated diseases. Therapies directed at control of the angiogenic processes could lead to the abrogation or mitigation of these diseases. Thus, angiogenesis is associated with any new tissue growth, whether normal or disease associated. Accordingly, the therapeutics described herein can be used to inhibit angiogenesis associated with any new tissue growth, whether normal or disease associated.

Therapeutics may be contacted with a tissue to prevent angiogenesis and/or cell proliferation, e.g., tumor growth, or any disease or disorder associated therewith. Diseases and processes that are mediated by angiogenesis include hemangioma, solid tumors, leukemia, metastasis, telangiectasia psoriasis scleroderma, pyogenic granuloma, myocardial angiogenesis, plaque neovascularization, coronary collaterals, cerebral collaterals, arteriovenous malformations, ischemic limb angiogenesis, corneal diseases, rubeosis, neovascular glaucoma, diabetic retinopathy, retrolental fibro-

plasia, arthritis, diabetic neovascularization, macular degeneration, wound healing, peptic ulcer, fractures, keloids, phemphigoid, trachoma, vasculogenesis, hematopoiesis, ovulation, menstruation, and placentation. The therapeutics described herein may also be used for treating or repressing the growth of a cancer or reducing tumor mass.

Angiogenesis is prominent in solid tumor formation and metastasis. Angiogenic factors have been found associated with several solid tumors such as rhabdomyosarcomas, retinoblastoma, Ewing sarcoma, neuroblastoma, and osteosarcoma. A tumor cannot expand without a blood supply to provide nutrients and remove cellular wastes. Tumors in which angiogenesis is important include solid tumors, and benign tumors such as acoustic neuroma, neurofibroma, trachoma and pyogenic granulomas. Prevention of angiogenesis could halt the growth of these tumors and the resultant damage to the animal due to the presence of the tumor.

Angiogenesis is important in two stages of tumor metastasis. The first stage where angiogenesis stimulation is important is in the vascularization of the tumor which allows tumor cells to enter the blood stream and to circulate throughout the body. After the tumor cells have left the primary site, and have settled into the secondary, metastasis site, angiogenesis must occur before the new tumor can grow and expand. Therefore, prevention of angiogenesis could lead to the prevention of metastasis of tumors and possibly contain the neoplastic growth at the primary site.

It should be noted that angiogenesis has been associated with blood-born tumors such as leukemias, any of various acute or chronic neoplastic diseases of the bone marrow in which unrestrained proliferation of white blood cells occurs, usually accompanied by anemia, impaired blood clotting, and enlargement of the lymph nodes, liver, and spleen. It is believed that angiogenesis plays a role in the abnormalities in the bone marrow that give rise to leukemia-like tumors.

The compositions described herein may be used for treating atherosclerosis. As such, compositions comprising the peptides described herein may be used to prevent or regress atherosclerosis growth or plaque formation.

Generally, the compositions described herein may be used for treating inflammatory-disorders, such as immune and non-immune inflammation, chronic articular rheumatism, psoriasis disorders associated with inappropriate or inopportune invasion of vessels, such as diabetic retinopathy, neovascular glaucoma, restenosis, capillary proliferation in atherosclerotic plaques and osteoporosis. Cancer-associated disorders that can be treated include solid tumors, solid tumor metastasis, angiofibromas, retrolental fibroplasia, hemangiomas, and Kaposi sarcomas.

One example of a disease mediated by angiogenesis is ocular neovascular disease. This disease is characterized by invasion of new blood vessels into the structures of the eye such as the retina or cornea. It is the most common cause of blindness and is involved in approximately twenty eye diseases. In age-related macular degeneration, the associated visual problems are caused by an ingrowth of chorioidal capillaries through defects in Bruch's membrane with proliferation of fibrovascular tissue beneath the retinal pigment epithelium. Angiogenic damage is also associated with diabetic retinopathy, retinopathy of prematurity, corneal graft rejection, neovascular glaucoma and retrolental fibroplasia.

Diseases associated with corneal neovascularization that can be treated as described herein include but are not limited to, diabetic retinopathy, retinopathy of prematurity, corneal graft rejection, neovascular glaucoma and retrolental fibroplasia, epidemic keratoconjunctivitis, Vitamin A deficiency, contact lens overwear, atopic keratitis, superior limbic kerati-

tis, pterygium keratitis sicca, sjogrens, acne rosacea, phlyctenulosis, syphilis, Mycobacteria infections, lipid degeneration, chemical burns, bacterial ulcers, fungal ulcers, Herpes simplex infections, Herpes zoster infections, protozoan infections, Kaposi sarcoma, Mooren ulcer, Terrien's marginal degeneration, mariginal keratolysis, trauma, rheumatoid arthritis, systemic lupus, polyarteritis, Wegeners sarcoidosis, Scieritis, Steven's Johnson disease, and periphigoid radial keratotomy.

Diseases associated with retinal/choroidal neovascularization that can be treated as described herein include, but are not limited to, diabetic retinopathy, macular degeneration, sickle cell anemia, sarcoid, syphilis, pseudoxanthoma elasticum, Pagets disease, vein occlusion, artery occlusion, carotid obstructive disease, chronic uveitis/vitritis, mycobacterial infections, Lyme's disease, systemic lupus erythematosus, retinopathy of prematurity, Eales disease, Bechets disease, infections causing a retinitis or choroiditis, presumed ocular histoplasmosis, Bests disease, myopia, optic pits, Stargarts disease, pars planitis, chronic retinal detachment, hyperviscosity syndromes, toxoplasmosis, trauma and post-laser complications. Other diseases include, but are not limited to, diseases associated with rubeosis (neovasculariation of the angle) and diseases caused by the abnormal proliferation of fibrovascular or fibrous tissue including all forms of proliferative vitreoretinopathy, whether or not associated with diabetes.

Another disease in which angiogenesis is believed to be involved is rheumatoid arthritis. The blood vessels in the synovial lining of the joints undergo angiogenesis. In addition to forming new vascular networks, the endothelial cells release factors and reactive oxygen species that lead to pannus growth and cartilage destruction. The factors involved in angiogenesis may actively contribute to, and help maintain, the chronically inflamed state of rheumatoid arthritis.

Factors associated with angiogenesis may also have a role in osteoarthritis. The activation of the chondrocytes by angiogenic-related factors contributes to the destruction of the joint. At a later stage, the angiogenic factors would promote new bone formation. Therapeutic intervention that prevents the bone destruction could halt the progress of the disease and provide relief for persons suffering with arthritis.

Chronic inflammation may also involve pathological angiogenesis. Such disease states as ulcerative colitis and Crohn's disease show histological changes with the ingrowth of new blood vessels into the inflamed tissues. Bartonellosis, a bacterial infection found in South America, can result in a chronic stage that is characterized by proliferation of vascular endothelial cells. Another pathological role associated with angiogenesis is found in atherosclerosis. The plaques formed within the lumen of blood vessels have been shown to have angiogenic stimulatory activity.

One of the most frequent angiogenic diseases of childhood is the hemangioma. In most cases, the tumors are benign and regress without intervention. In more severe cases, the tumors progress to large cavernous and infiltrative forms and create clinical complications. Systemic forms of hemangiomas, the hemangiomatoses, have a high mortality rate. Therapy-resistant hemangiomas exist that cannot be treated with therapeutics currently in use.

Angiogenesis is also responsible for damage found in hereditary diseases such as Osler-Weber-Rendu disease, or hereditary hemorrhagic telangiectasia. This is an inherited disease characterized by multiple small angiomas, tumors of blood or lymph vessels. The angiomas are found in the skin and mucous membranes, often accompanied by epistaxis

(nosebleeds) or gastrointestinal bleeding and sometimes with pulmonary or hepatic arteriovenous fistula.

Another disease that can be treated according to the present invention is acquired immune deficiency syndrome.

The therapeutic peptides may also be used for decreasing overproliferation of normal, vascularized tissues, such as adipose tissue, benign polyps, hypertrophied cardiac tissue, hypertrophied renal tissue, hypertrophied prostatic tissue, tissue containing amyloid deposits, and uterine fibroids. The therapeutics may be administered in an amount effective to reduce the vascular supply to the tissue, or to decrease the size or growth of the vascularized tissue, such as adipose tissue, polyps (e.g., intestinal or nose polyps), and muscle (including cardiac) tissue. Accordingly, subject that may be treated include a subject having polypsis, and enlarged prostate, cardiac or renal hypertrophy or is obese or overweight. The peptide therapeutics may be administered in an amount and time period which results in blood levels regulating the size and/or growth of the vascularized tissue to be treated.

The therapies for losing weight are applicable to both normal overweight individuals and individuals with genetic defects. The method should also be useful in most cases involving weight gains due to hormonal or metabolic defects or drug side effects. In addition to promoting loss of body fat while maintaining lean body mass and being able to sustain weight loss during chronic administration, other benefits of the treatments include normalization of blood glucose levels in obesity related diabetes.

In addition, any disease or secondary condition developing as a result from a disease associated with angiogenesis, which can be treated according to the methods described herein, may also be treated. For example, any condition resulting from being overweight or obese may be treated or prevented as described herein. Exemplary diseases include hyperlipidemia, dyslipogenesis, hypercholesterolemia, impaired glucose tolerance, high blood glucose sugar level, syndrome X, hypertension, atherosclerosis and lipodystrophy, hypertension, high blood cholesterol, dyslipidemia, type 2 diabetes, insulin resistance, glucose intolerance, hyperinsulinemia, coronary heart disease, angina pectoris, congestive heart failure, stroke, gallstones, cholecystitis and cholelithiasis, gout, osteoarthritis, obstructive sleep apnea and respiratory problems, some types of cancer (such as endometrial, breast, prostate, colon, lung, liver, lymph node, kidney, pancreas, ovary, spleen, small intestine, stomach, skin, testes, head and neck, esophagus, brain (glioblastomas, medulloblastoma, astrocytoma, oligodendroglioma, ependymomas), blood cells, bone marrow or other tissue), complications of pregnancy, poor female reproductive health (such as menstrual irregularities, infertility, irregular ovulation), bladder control problems (such as stress incontinence); uric acid nephrolithiasis; and psychological disorders (such as depression, eating disorders, distorted body image, and low self esteem).

Angiogenesis is also involved in normal physiological processes such as reproduction and wound healing. Angiogenesis is an important step in ovulation and also in implantation of the blastula after fertilization. Prevention of angiogenesis could be used to induce amenorrhea, to block ovulation or to prevent implantation by the blastula.

In wound healing, excessive repair or fibroplasia can be a detrimental side effect of surgical procedures and may be caused or exacerbated by angiogenesis. Adhesions are a frequent complication of surgery and lead to problems such as small bowel obstruction. Therefore in this and in other situations, it may be desirable to prevent wound healing with the therapeutic peptides described herein.

Peptides may be provided in pharmaceutical compositions and administered to a subject in need thereof. Peptides may also be contacted with a tissue or cells in vitro.

Therapeutic methods generally comprise administering to a subject, e.g., a subject in need thereof, a therapeutically effective amount of a therapeutic, e.g., a peptide or variant or derivative thereof or nucleic acid encoding such, as, e.g., described herein. Subjects may be mammals, such as humans, non-human primates, canines, felines, equines, porcines, ovines, bovines, sheep, mice and rat. In particular, methods described herein can be used for veterinary purposes. The method may comprise first diagnosing a subject with a disease or disorder in which administration of a peptide described herein may be beneficial, such as a malignant or benign tumor growth. The method may also comprise determining the effect of the therapeutic a certain time after its administration. For example, the size of the tumor growth may be evaluated about one week, one month or two months after starting of the treatment. The evaluation may also comprising obtaining a sample of tissue, e.g., a tumor, and determining the level of angiogenesis.

The therapeutics can be administered in a "growth inhibitory amount," i.e., an amount of the peptide that is therapeutically effective to inhibit or decrease proliferation of cells or tissues. The therapeutics may also be administered in an "antiangiogenic amount," i.e., an amount of a therapeutic that is therapeutically effective to inhibit or decrease angiogenesis. The therapeutics may be administered to mammals, preferably humans, either alone or, in combination with pharmaceutically acceptable carriers, excipients or diluents, in a pharmaceutical composition, according to standard pharmaceutical practice. Therapeutics may be administered directly into the tissue in which one desires to inhibit angiogenesis or tumor growth. The therapeutics may also be administered orally or parenterally, including intravenously, intramuscularly, intraperitoneally, subcutaneously, rectally and topically. One or more therapeutics can be injected directly into a tumor of the subject to be treated.

The peptide or variant or derivative thereof may be co-administered with zinc. For example a therapeutic composition may be administered in a therapeutically effective dose (see below) together with a therapeutically effective dose of Zn^{2+} (see below). The peptide or variant or derivative thereof may be combined with the zinc solution prior to administration or at the time of administration, such as to allow zinc to interact with the peptide. Alternatively, the peptide or variant or derivative thereof is administered separately from the zinc, either before or after administration of the zinc, provided that both are present in the circulation at least during a common period. For example, a solution of zinc may be administered about a few minutes to a few hours before or after administration of the peptide. In yet other embodiments, no zinc is administered to a subject to whom a peptide is administered. Peptides that are administered may, however, still bind zinc when the subject has zinc in their blood circulation.

Toxicity and therapeutic efficacy of the therapeutics can be determined by standard pharmaceutical procedures in cell cultures or experimental animals, e.g., for determining the LD_{50} (the dose lethal to 50% of the population) and the ED_{50} (the dose therapeutically effective in 50% of the population). The dose ratio between toxic and therapeutic effects is the therapeutic index and it can be expressed as the ratio LD_{50}/ED_{50} . Reagents which exhibit large therapeutic indices are preferred. While reagents that exhibit toxic side effects may be used, care should be taken to design a delivery system that

targets such reagents to the site of affected tissue in order to, e.g., minimize potential damage to normal cells and, thereby, reduce side effects.

The data obtained from cell culture assays and animal studies can be used in formulating a range of dosage for use in humans. The dosage of such therapeutics lies preferably within a range of circulating concentrations that include the ED₅₀ with little or no toxicity. The dosage may vary within this range depending upon the dosage form employed and the route of administration utilized. For any therapeutic used, the therapeutically effective dose can be estimated initially from cell culture assays. A dose may be formulated in animal models to achieve a circulating plasma concentration range that includes the IC₅₀ (i.e., the concentration of the test therapeutic which achieves a half-maximal inhibition of symptoms) as determined in cell culture. Such information can be used to more accurately determine useful doses in humans.

The dosage of the therapeutic will depend on the disease state or condition being treated and other clinical factors such as weight and condition of the human or animal and the route of administration of the compound. For treating humans or animals, between approximately 0.5 mg/kilogram to 500 mg/kilogram of the therapeutic can be administered. A more preferable range is about 1 mg/kilogram to about 100 mg/kilogram or from about 2 mg/kilogram to about 50 mg/kilogram with the most preferable range being from about 2 mg/kilogram to about 10 mg/kilogram. Depending upon the half-life of the therapeutic in the particular animal or human, the therapeutic can be administered between several times per day to once a week. It is to be understood that the methods have application for both human and veterinary use. The methods of the present invention contemplate single as well as multiple administrations, given either simultaneously or over an extended period of time.

When the peptide or analog or derivative thereof is administered with zinc, zinc may be included in the therapeutic composition comprising the peptide or analog or derivative thereof in a concentration from about 0.1 to about 100 mg/kg/day; about 1 to about 10 mg/kg/day; or about 2-5 mg/kg/day. Zinc may be administered in the form of Zn²⁺ or a salt thereof. The amount of zinc may vary depending on the amount of zinc in a subject's circulation (e.g., blood) and on the amount of peptide administered to the subject. The amount of zinc that may be necessary can be determined, e.g., by taking a blood sample of a subject having received a particular dose of peptide alone or with zinc and determining the amount of peptide to which zinc is complexed, e.g., as described above.

Pharmaceutical compositions containing a therapeutic may be in a form suitable for oral use, for example, as tablets, troches, lozenges, aqueous or oily suspensions, dispersible powders or granules, emulsions, hard or soft capsules, or syrups or elixirs. Compositions intended for oral use may be prepared according to any method known to the art for the manufacture of pharmaceutical compositions and such compositions may contain one or more agents selected from the group consisting of sweetening agents, flavoring agents, coloring agents and preserving agents in order to provide pharmaceutically elegant and palatable preparations. Tablets may contain the active ingredient (i.e., therapeutic) in admixture with non-toxic pharmaceutically acceptable excipients which are suitable for the manufacture of tablets. These excipients may be for example, inert diluents, such as calcium carbonate, sodium carbonate, lactose, calcium phosphate or sodium phosphate; granulating and disintegrating agents, for example, microcrystalline cellulose, sodium crosscarmellose, corn starch, or alginic acid; binding agents, for example starch, gelatin, polyvinyl-pyrrolidone or acacia, and lubricat-

ing agents, for example, magnesium stearate, stearic acid or talc. The tablets may be uncoated or they may be coated by known techniques to mask the unpleasant taste of the drug or delay disintegration and absorption in the gastrointestinal tract and thereby provide a sustained action over a longer period. For example, a water soluble taste masking material such as hydroxypropylmethyl-cellulose or hydroxypropyl-cellulose, or a time delay material such as ethyl cellulose, cellulose acetate butyrate may be employed.

Formulations for oral use may also be presented as hard gelatin capsules wherein the active ingredient is mixed with an inert solid diluent, for example, calcium carbonate, calcium phosphate or kaolin, or as soft gelatin capsules wherein the active ingredient is mixed with water soluble carrier such as polyethyleneglycol or an oil medium, for example peanut oil, liquid paraffin, or olive oil.

Aqueous suspensions may contain the active material in admixture with excipients suitable for the manufacture of aqueous suspensions. Such excipients are suspending agents, for example sodium carboxymethylcellulose, methylcellulose, hydroxypropylmethyl-cellulose, sodium alginate, polyvinyl-pyrrolidone, gum tragacanth and gum acacia; dispersing or wetting agents may be a naturally-occurring phosphatide, for example lecithin, or condensation products of an alkylene oxide with fatty acids, for example polyoxyethylene stearate, or condensation products of ethylene oxide with long chain aliphatic alcohols, for example heptadecaethylene-oxycetanol, or condensation products of ethylene oxide with partial esters derived from fatty acids and a hexitol such as polyoxyethylene sorbitol monooleate, or condensation products of ethylene oxide with partial esters derived from fatty acids and hexitol anhydrides, for example polyethylene sorbitan monooleate. The aqueous suspensions may also contain one or more preservatives, for example ethyl, or n-propyl p-hydroxybenzoate, one or more coloring agents, one or more flavoring agents, and one or more sweetening agents, such as sucrose, saccharin or aspartame.

Oily suspensions may be formulated by suspending the active ingredient in a vegetable oil, for example arachis oil, olive oil, sesame oil or coconut oil, or in mineral oil such as liquid paraffin. The oily suspensions may contain a thickening agent, for example beeswax, hard paraffin or cetyl alcohol. Sweetening agents such as those set forth above, and flavoring agents may be added to provide a palatable oral preparation. These compositions may be preserved by the addition of an anti-oxidant such as butylated hydroxyanisole or alpha-tocopherol.

Dispersible powders and granules suitable for preparation of an aqueous suspension by the addition of water provide the active ingredient in admixture with a dispersing or wetting agent, suspending agent and one or more preservatives. Suitable dispersing or wetting agents and suspending agents are exemplified by those already mentioned above. Additional excipients, for example sweetening, flavoring and coloring agents, may also be present. These compositions may be preserved by the addition of an anti-oxidant such as ascorbic acid.

Pharmaceutical compositions may also be in the form of an oil-in-water emulsions. The oily phase may be a vegetable oil, for example olive oil or arachis oil, or a mineral oil, for example liquid paraffin or mixtures of these. Suitable emulsifying agents may be naturally-occurring phosphatides, for example soy bean lecithin, and esters or partial esters derived from fatty acids and hexitol anhydrides, for example sorbitan monooleate, and condensation products of the said partial esters with ethylene oxide, for example polyoxyethylene sor-

bitan monooleate. The emulsions may also contain sweetening, flavouring agents, preservatives and antioxidants.

Syrups and elixirs may be formulated with sweetening agents, for example glycerol, propylene glycol, sorbitol or sucrose. Such formulations may also contain a demulcent, a preservative, flavoring and coloring agents and antioxidant.

The pharmaceutical compositions may be in the form of a sterile injectable aqueous solution. Among the acceptable vehicles and solvents that may be employed are water, Ringier's solution and isotonic sodium chloride solution.

The sterile injectable preparation may also be a sterile injectable oil-in-water microemulsion where the active ingredient is dissolved in the oily phase. For example, the active ingredient may be first dissolved in a mixture of soybean oil and lecithin. The oil solution then introduced into a water and glycerol mixture and processed to form a microemulsion.

The injectable solutions or microemulsions may be introduced into a patient's blood-stream by local bolus injection. Alternatively, it may be advantageous to administer the solution or microemulsion in such a way as to maintain a constant circulating concentration of the instant compound. In order to maintain such a constant concentration, a continuous intravenous delivery device may be utilized. An example of such a device is the Deltec CADD-PLUS™ model 5400 intravenous pump. For example, one could generate blood levels of therapeutic peptides in the range of about 100 to 500 ng/ml, or about 200 to 400 ng/ml or about 250-300 ng/ml.

The pharmaceutical compositions may be in the form of a sterile injectable aqueous or oleagenous suspension for intramuscular and subcutaneous administration. This suspension may be formulated according to the known art using those suitable dispersing or wetting agents and suspending agents which have been mentioned above. The sterile injectable preparation may also be a sterile injectable solution or suspension in a non-toxic parenterally-acceptable diluent or solvent, for example as a solution in 1,3-butane-diol. In addition, sterile, fixed oils are conventionally employed as a solvent or suspending medium. For this purpose any bland fixed oil may be employed including synthetic mono- or diglycerides. In addition, fatty acids such as oleic acid find use in the preparation of injectables.

In certain embodiments, it may be preferable to administer a therapeutic locally, such as by local injection. For example, a therapeutic may be injected directly into the tissue showing excessive proliferation in which one desired to inhibit angiogenesis. In one embodiment, a therapeutic is administered locally into a tumor bed.

In one embodiment, a therapeutic peptide is incorporated into a topical formulation containing, e.g., a topical carrier that is generally suited to topical drug administration and comprising any such material known in the art. The topical carrier may be selected so as to provide the composition in the desired form, e.g., as an ointment, lotion, cream, microemulsion, gel, oil, solution, or the like, and may be comprised of a material of either naturally occurring or synthetic origin. It is preferable that the selected carrier not adversely affect the active agent or other components of the topical formulation. Examples of suitable topical carriers for use herein include water, alcohols and other nontoxic organic solvents, glycerin, mineral oil, silicone, petroleum jelly, lanolin, fatty acids, vegetable oils, parabens, waxes, and the like. Formulations may be colorless, odorless ointments, lotions, creams, microemulsions and gels.

Various additives, known to those skilled in the art, may be included in formulations, e.g., topical formulations. Examples of additives include, but are not limited to, solubilizers, skin permeation enhancers, opacifiers, preservatives

(e.g., anti-oxidants), gelling agents, buffering agents, surfactants (particularly nonionic and amphoteric surfactants), emulsifiers, emollients, thickening agents, stabilizers, humectants, colorants, fragrance, and the like. Inclusion of solubilizers and/or skin permeation enhancers is particularly preferred, along with emulsifiers, emollients and preservatives. An optimum topical formulation comprises approximately: 2 wt. % to 60 wt. %, preferably 2 wt. % to 50 wt. %, solubilizer and/or skin permeation enhancer; 2 wt. % to 50 wt. %, preferably 2 wt. % to 20 wt. %, emulsifiers; 2 wt. % to 20 wt. % emollient; and 0.01 to 0.2 wt. % preservative, with the active agent and carrier (e.g., water) making of the remainder of the formulation.

Other active agents may also be included in formulations, e.g., other anti-inflammatory agents, analgesics, antimicrobial agents, antifungal agents, antibiotics, vitamins, antioxidants, and sunblock agents commonly found in sunscreen formulations including, but not limited to, anthranilates, benzophenones (particularly benzophenone-3), camphor derivatives, cinnamates (e.g., octyl methoxycinnamate), dibenzoyl methanes (e.g., butyl methoxydibenzoyl methane), p-aminobenzoic acid (PABA) and derivatives thereof, and salicylates (e.g., octyl salicylate).

Topical formulations may also be used as preventive, e.g., chemopreventive, compositions. When used in a chemopreventive method, susceptible skin is treated prior to any visible condition in a particular individual.

Therapeutics may also be administered in the form of a suppository for rectal administration of the drug. These compositions can be prepared by mixing the drug with a suitable non-irritating excipient which is solid at ordinary temperatures but liquid at the rectal temperature and will therefore melt in the rectum to release the drug. Such materials include cocoa butter, glycerinated gelatin, hydrogenated vegetable oils, mixtures of polyethylene glycols of various molecular weights and fatty acid esters of polyethylene glycol.

For topical use, creams, ointments, jellies, solutions or suspensions, etc., containing the therapeutics may be employed. For purposes of this application, topical application shall include mouth washes and gargles.

Therapeutics may be administered in intranasal form via topical use of suitable intranasal vehicles and delivery devices, or via transdermal routes, using those forms of transdermal skin patches well known to those of ordinary skill in the art. To be administered in the form of a transdermal delivery system, the dosage administration will, of course, be continuous rather than intermittent throughout the dosage regimen.

The therapeutic peptides may be administered on a dose-schedule that maintains a constant concentration in the circulation. They may also be administered on a dose-schedule in which therapy is periodically discontinued, e.g., a once daily bolus injection.

Single therapeutic peptides or variants or derivatives thereof may be administered. Alternatively, two or more different peptides may be co-administered. For example, a therapeutic peptide may be administered on its own or it may be co-administered with other angiogenic inhibitors, such as TNP-470, described by U.S. Pat. No. 5,290,807; Angiostatin, described by U.S. Pat. No. 5,639,725; Endostatin, and Thalidomide. Other angiogenic inhibitors are described in Genetic Engineering News, Oct. 1, 1998 and in U.S. Pat. No. 6,306,819.

In other embodiments, the therapeutics are co-administered with other well known therapeutic agents that are selected for their particular usefulness against the condition that is being treated. For example, the instant therapeutics

may be useful in combination with known anti-cancer and cytotoxic agents. Similarly, the instant therapeutics may be useful in combination with agents that are effective in the treatment and prevention of benign or malignant tumors. The therapeutic agents may be added to chemotherapy, radiotherapy or used in combination with immunotherapy or vaccine therapy.

Drugs that can be co-administered to a subject being treated with a therapeutic described herein include antineoplastic agents selected from vinca alkaloids, epipodophyllotoxins, anthracycline antibiotics, actinomycin D, plicamycin, puromycin, gramicidin D, taxol, colchicine, cytochalasin B, emetine, maytansine, or amsacrine.

Classes of compounds that can be used as the chemotherapeutic agent (antineoplastic agent) include: alkylating agents, antimetabolites, natural products and their derivatives, hormones and steroids (including synthetic analogs), and synthetics. Examples of compounds within these classes are given below. Alkylating agents (including nitrogen mustards, ethylenimine derivatives, alkyl sulfonates, nitrosoureas and triazenes): Uracil mustard, Chloromethine, Cyclophosphamide (CytosanTM), Ifosfamide, Melphalan, Chlorambucil, Pipobroman, Triethylene-melamine, Triethylenethiophosphoramine, Busulfan, Carmustine, Lomustine, Streptozocin, Dacarbazine, and Temozolomide. Antimetabolites (including folic acid antagonists, pyrimidine analogs, purine analogs and adenosine deaminase inhibitors): Methotrexate, 5-Fluorouracil, Floxuridine, Cytarabine, 6-Mercaptopurine, 6-Thioguanine, Fludarabine phosphate, Pentostatin, and Gemcitabine. Natural products and their derivatives (including vinca alkaloids, antitumor antibiotics, enzymes, lymphokines and epipodophyllotoxins): Vinblastine, Vincristine, Vindesine, Bleomycin, Dactinomycin, Daunorubicin, Doxorubicin, Epirubicin, Idarubicin, paclitaxel (paclitaxel is commercially available as Taxol^{RTM} and is described in more detail below in the subsection entitled "Microtubule Affecting Agents"), Mithramycin, Deoxycoformycin, Mitomycin-C, L-Asparaginase, Interferons (especially IFN- α), Etoposide, and Teniposide. Hormones and steroids (including synthetic analogs): 17.alpha.-Ethinylestradiol, Diethylstilbestrol, Testosterone, Prednisone, Fluoxymesterone, Dromostanolone propionate, Testolactone, Megestrolacetate, Tamoxifen, Methylprednisolone, Methyltestosterone, Prednisolone, Triamcinolone, Chlorotrianisene, Hydroxyprogesterone, Aminoglutethimide, Estramustine, Medroxyprogesteroneacetate, Leuprolide, Flutamide, Toremifene, Zoladex. Synthetics (including inorganic complexes such as platinum coordination complexes): Cisplatin, Carboplatin, Hydroxyurea, Amsacrine, Procarbazine, Mitotane, Mitoxantrone, Levamisole, and Hexamethylmelamine.

Methods for the safe and effective administration of most of these chemotherapeutic agents are known to those skilled in the art. In addition, their administration is described in the standard literature. For example, the administration of many of the chemotherapeutic agents is described in the "Physicians' Desk Reference" (PDR), e.g., 2004 edition (Thomson PDR, Montvale, N.J. 07645-1742, USA).

If formulated as a fixed dose, such combination products employ the combinations described herein within the dosage range described herein and the other pharmaceutically active agent(s) within its approved dosage range. Combinations of the instant invention may also be used sequentially with known pharmaceutically acceptable agent(s) when a multiple combination formulation is inappropriate.

Radiation therapy, including x-rays or gamma rays which are delivered from either an externally applied beam or by

implantation of tiny radioactive sources, may also be used in combination with a therapeutic described herein to treat cancer.

When a therapeutic is administered into a human subject, the daily dosage will normally be determined by the prescribing physician with the dosage generally varying according to the age, weight, and response of the individual patient, as well as the severity of the patient's symptoms.

Therapeutic peptides or analogs thereof described herein can be labeled isotopically or with other molecules or proteins for use in the detection and visualization of endostatin binding sites with state of the art techniques, including, but not limited to, positron emission tomography, autoradiography, flow cytometry, radioreceptor binding assays, and immunohistochemistry. Labeled peptides or analogs thereof may be used for detecting and quantifying the presence of an antibody specific for an endostatin in a body fluid.

A person of skill in the art will understand that nucleic acids encoding endostatin peptides can be used instead of the peptides in most embodiments described herein. For example, angiogenesis can be inhibited in a tissue by introducing into the tissue a nucleic acid encoding a peptide described herein. A nucleic acid may be linked to or comprise a transcriptional control element, such as a promoter or an enhancer. A nucleic acid may further be included in a vector, such as an expression vector. Exemplary expression vectors include viral vectors, such as an adenovirus and an adeno associated virus (AAV) vector. When using a non-viral vector, various methods can be used for facilitating entry of the nucleic acid into a cell, such as liposomes.

Exemplary Kits

Also provided herein are kits, e.g., therapeutic kits. A kit may comprise a therapeutic peptide described herein and optionally a device for administration of the therapeutic peptide. A kit may also comprise a therapeutic peptide in a lyophilized form and a solution or buffer to solubilize the therapeutic peptide. A kit may also comprise instructions for use.

The practice of the present invention will employ, unless otherwise indicated, conventional techniques of cell biology, cell culture, molecular biology, transgenic biology, microbiology, recombinant DNA, and immunology, which are within the skill of the art. Such techniques are explained fully in the literature. See, for example, *Molecular Cloning A Laboratory Manual*, 2nd Ed., ed. by Sambrook, Fritsch and Maniatis (Cold Spring Harbor Laboratory Press: 1989); *DNA Cloning*, Volumes I and II (D. N. Glover ed., 1985); *Oligonucleotide Synthesis* (M. J. Gait ed., 1984); Mullis et al. U.S. Pat. No. 4,683,195; *Nucleic Acid Hybridization* (B. D. Hames & S. J. Higgins eds. 1984); *Transcription And Translation* (B. D. Hames & S. J. Higgins eds. 1984); *Culture Of Animal Cells* (R. I. Freshney, Alan R. Liss, Inc., 1987); *Immobilized Cells And Enzymes* (IRL Press, 1986); B. Perbal, *A Practical Guide To Molecular Cloning* (1984); the treatise, *Methods In Enzymology* (Academic Press, Inc., N.Y.); *Gene Transfer Vectors For Mammalian Cells* (J. H. Miller and M. P. Calos eds., 1987, Cold Spring Harbor Laboratory); *Methods In Enzymology*, Vols. 154 and 155 (Wu et al. eds.), *Immunochemical Methods In Cell And Molecular Biology* (Mayer and Walker, eds., Academic Press, London, 1987); *Handbook Of Experimental Immunology*, Volumes I-IV (D. M. Weir and C. C. Blackwell, eds., 1986); *Manipulating the Mouse Embryo*, (Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y., 1986).

Exemplification

The invention now being more generally described, it will now be more readily understood by reference to the following

examples, which are included merely for purposes of illustration of certain aspects and embodiments of the present invention, and are not intended to limit the invention.

EXAMPLE 1

Identification of a 27 Amino Acid Endostatin Peptide that is Responsible for Anti-Tumor Activity

Overlapping peptides with 24-27 amino acids derived from both mouse endostatin and human endostatin were synthesized (Table 1).

TABLE 1

Overlapping mouse and human endostatin peptides			
Name	Sequence		
Mouse Peptides			
mP1:	HTHQDFQPVLHLVALNTPLSGGMRGIR;	(SEQ ID NO: 4)	
mP2:	MRGIRGA DFQAFQQARA VGLSGTFR;	(SEQ ID NO: 136)	
mP3:	TFRAFLSSRLQDLYSIVRRADRGSV;	(SEQ ID NO: 137)	
mP4:	GSVPIVNLKDEVLSPSWDSLFGSGSQ;	(SEQ ID NO: 138)	
mP5:	GSQGQVQPGARIFSFDRDVLRLHQA;	(SEQ ID NO: 139)	
mP6:	HPAWPQKSVWHGSDPSGRRLMESY;	(SEQ ID NO: 140)	
mP7:	ETWRTETTGATGQASSLLSGRLLEQ;	(SEQ ID NO: 141)	
mP8:	KAASAHNSYIVLAIENSFMTSFSKKK.	(SEQ ID NO: 142)	
Human Peptides			
hP1	1 HSHRDFQPVLHLVALNSPLSGGMRG	25	(SEQ ID NO: 6)
hP2	23 MRGIRGADFQAFQQARAVGLAGTFR	47	(SEQ ID NO: 143)
hP3	45 TFRAFLSSRLQDLYSIVRRADRAAV	69	(SEQ ID NO: 144)
hP4	67 AAVPIVNLKDELLFPSWEALFGSGE	91	(SEQ ID NO: 145)
hP5	89 GSEGPLKPGARIFSFDRDVLRLHPT	113	(SEQ ID NO: 146)
hP6	111 HPTWPQKSVWHGSDPNRRRLTESY	134	(SEQ ID NO: 147)
hP7	136 ETWRTAPSATGQASSLLGGRLLGQ	160	(SEQ ID NO: 148)
hP8	158 LGQSAASAHAYIVLAIENSFMTASKKK	183	(SEQ ID NO: 149)

Peptides were approximately 1/7th-1/8th the size of full length endostatin. Three cysteines, 33,165,173, were replaced by alanines (underlined in Table 1), and cysteine135 was omitted, to prevent the formation of disulfide bonds. Two additional lysines were added at the C-terminal of hP8 to increase its solubility (double underlined). Most peptides were water soluble, except for hP2 at high concentrations (>2.5 mg/ml). Also, all peptides were approximately 70% pure. However, no difference in tumor inhibition was observed when peptides of more than 95% purity were used.

These peptides were initially tested for anti-tumor activity employing the human pancreatic tumor cells BxPC-3 cells, which were implanted in the s.c. dorsa of SCID mice. For systemic treatment, human endostatin peptides, P1-P8, were administered subcutaneously (s.c.) twice a day at 7 mg/kg/d due to the high clearance rate from the mouse circulation. Full-length Fc-Endostatin, hFcES was administered subcutaneously only once per day at a dose of 20 mg/kg/d. PBS was used as a control. Tumors were measured every three days and

the final measurements at 28 days are shown in FIG. 1 (T/C is indicated in each bar and the group sizes had an n equal to 3).

The N-terminal hP1 peptide of endostatin inhibited BxPC-3 by 39% (p=0.077) and full-length endostatin by 44% (p=0.0057) (FIG. 1). Two other peptides, hP2 and hP5, also showed some small anti-tumor activity, where hP2 inhibited BxPC-3 by 19% (p=0.48) and hP5 by 29% (p=0.15). Other peptides had no effect (FIG. 1). Thus, most of the anti-tumor activity was associated with the N-terminal hP1 peptide compared to full-length endostatin. Although tumor inhibition by hP1, hP2 and hP5 was not statistically significant due to the

small number of mice per group (n=3), the trends suggesting tumor inhibition by these peptides prompted further study of such peptides on tumor inhibition in the murine LLC model. Moreover, peptides and full-length endostatin were not at equimolar concentrations. However, this data suggests that the anti-tumor property of endostatin may be located within its N-terminal domain.

The N-terminal 27 Amino Acid Peptide of Endostatin is Responsible for Its Anti-Tumor Property

The murine LLC tumor model (O'Reilly et al. (1994) Cell, 79: 315-328) was used to further characterize these peptides because this tumor grows more rapidly than BxPC-3 cells, allowing for shorter treatment periods. Murine analogs of the endostatin peptides were synthesized. The only difference between the human and murine peptides was that murine P1 peptide contained 27 amino acids instead of 25 amino acids for human P1. We tested only those peptides that showed any anti-tumor activity in the previous experiment (see FIG. 1).

Unlike the BxPC-3 treatment, LLC tumors were treated at equimolar concentrations of murine endostatin and murine peptides (mP1, mP2, mP5, and mP6). mP6 was used as a representative control peptide, because it showed no anti-tumor activity (see FIG. 1). Control mice were treated with PBS. In these experiments, LLC cells were implanted into the s.c. dorsa of C57B1/6J mice, and systemically treated. Peptides (mP1, mP2, mP5, and mP6) were injected (s.c.) twice a day at a dose of 2.8 mg/kg/d, whereas endostatin and PBS were administered once a day at a dose of 20 mg/kg/d. The N-terminal mP1 endostatin peptide inhibited LLC by 44% ($p < 0.035$), which is comparable to the inhibition by full-length endostatin (53%, $p < 0.01$) (FIG. 2A; T/C is indicated in the figure). No or insignificant activity was detected employing mP2, mP5 and mP6 at the same concentration as the mP1 peptide (FIG. 2A). Thus, this result suggests that the 27 amino acids mP1 peptide contains all of the anti-tumor activity associated with endostatin.

To determine the effect on angiogenesis after mP1 treatment, LLC tumors treated with mP1, mP2, and endostatin at equimolar concentration were analyzed for vessel density (CD31). FIG. 2B shows CD31 staining of LLC sections from mice treated with PBS (control), Fc-Endostatin (20 mg/kg/d), mP1 (2.8 mg/kg/d), and mP2 (2.8 mg/kg/d). In each case, peptides were administered twice a day s.c. and LLC tumor sections from day 13 were formalin-fixed paraffin-embedded and then stained with CD31 (PECAM). FIG. 2C show the determination of vessel density and the Y axis is expressed as % CD31/high power field (hpf). Treatment of LLC with mP1 and endostatin reduced vessel density significantly (approximately 65%, $p < 0.015$), while mP2 and PBS had no effect. Treatment with mP5 and mP6 showed similar results as mP2 treatment. These results suggest that mP1 can inhibit LLC tumor growth by reducing vessel density in similar manner to full length endostatin.

Histidines at Position 1 and 3 of Endostatin are Critical for Zinc Binding

The crystal structure of endostatin reveals a highly folded molecule (FIG. 2D). However, the N-terminal region resembles a random coil structure consistent with our analysis that a synthetic peptide corresponding to this domain can mimic the native molecule (FIG. 2D).

Because the endostatin N-terminal domain is responsible for its anti-tumor activity, we wanted to investigate the mP1 peptide further. There is an atom of Zinc (Zn) associated with each molecule of endostatin (Ding et al. (1998) Proc Natl Acad Sci USA, 95: 10443-10448). Based on our crystal structure analysis, three histidines at positions 1, 3, and 11 plus, aspartic at position 76, form the four coordinates for this Zn atom (Ding et al. (1998) Proc Natl Acad Sci USA, 95: 10443-10448). mP1 contains the three histidines mentioned above. This raises the possibility that this peptide is capable of binding Zn, by having a molecule of water occupying the fourth coordinate (FIG. 3A, left panel). It has been shown previously that mutations of histidines 1 and 3 disrupt Zn binding of endostatin (Boehm et al. (1998) Biochem Biophys Res Commun, 252: 190-194). Therefore, a mutant of peptide mP1 was synthesized where the histidines at position 1 and 3 were mutated to alanines. This mutant peptide was called mP1-H1/3A (also referred to mP1-H) and has the following amino acid sequence: ATAQDFQPVLHLVALNTPLSGGMRGIR (SEQ ID NO: 150). The sequences of mP1 and mP1-H1/3A are also indicated in FIG. 3A.

To determine the Zn binding capacity of mP1 and mP1-H1/3A flame atomic absorption was performed. Each peptide was dissolved in 20 mM Tris, pH 8.0 at a concentration of 0.5

mg/ml, mixed with excess Zn chloride (1 mM), and extensively dialyzed against the above buffer for 72 hours with three changes in dialysis solution (molecular weight cut off (MWCO)=1000 kDa; the molecular weight of the peptides was taken to be 3000 kDa). Atomic absorption readings of the final zinc concentrations ($\mu\text{g/ml}$) were determined to be 9.63 and 1.05 for mP1 and mP1-H1/3A, respectively. These data yielded zinc ratios of 0.1 per molecule of mP1-H1/3A and 0.9 for mP1 (FIG. 3B). Therefore, mutating the histidines at position 1 and 3 to alanines abolished Zn binding (FIG. 3A, right panel).

The Zinc Binding Domain of Endostatin is Important for Its Anti-Tumor Activity

To determine if Zn binding is also important for the anti-tumor property of endostatin, mP1 and mP1-H1/3A were tested using the LLC tumor model. Peptides were administered twice a day (s.c.) at a dose of 2.8 mg/kg/d. Peptide mP1 inhibited LLC by 42% ($p = 0.031$), whereas mP1-H1/3A had no effect (FIG. 3C). To determine if there was a difference in angiogenesis, vessel density (CD31) of LLC tumors was analyzed after mP1 and mP1-H1/3A treatment. There was a considerable decrease in vessel density after mP1 treatment (67% reduction, $p < 0.01$), whereas mP1-H1/3A was similar to PBS (FIGS. 3D and 3E). These data suggest that Zn binding is important for the anti-tumor property of endostatin. The unpaired Student t test was used for statistical analysis.

Additional treatments included subcutaneously injection the mP1-H peptide together with 1 mM of Zn^{2+} ; subcutaneously injecting the mP1 peptide alone or with 1 mM of Zn^{2+} ; or subcutaneously injecting PBS (FIG. 4). The results, which are shown in FIGS. 3 and 4, indicated that tumor volume was essentially not affected by the administration of the mP1-H1/3A peptide, in which histidines 1 and 3 are changed to alanines, since the tumor mass was similar to that obtained when injecting PBS (the negative control).

Thus, these results indicate that the histidines at positions 1 and 3 in N-terminal endostatin peptides cannot be replaced by alanines and that their substitution with another amino acid would likely also reduce or eliminate the anti-tumor effect of the peptides. The results further indicate that it may be beneficial to include zinc in the composition comprising the peptide, unless there is zinc already present in the circulation.

The N-Terminal Fragment of Endostatin Retains Its Capacity to Inhibit Endothelial Cell Migration

Endostatin peptides were tested for anti-endothelial cell migration activity. Inhibition of VEGF-induced migration of human microvascular endothelial cells (HMVECs) was determined using several doses of endostatin peptides (FIG. 5). Human peptides were used because the cells were of human origin and the migration response of the HMVECs was assayed using a modified Boyden chamber. VEGF (5 ng/ml) was used a chemotactic agent and cells were challenged with human endostatin (Entremed; EM-ES), human Fc-Endostatin (nFcES), human P1 (hP1), human P2 (hP2), human P6 (hP6), and human P1-H1/3A (hP1-H1/3A). Total migration per membrane was quantified from the capture images using Scion Image Software (National Institutes of Health) and the unpaired Student T test was used for statistical analysis.

Human recombinant endostatin (Entremed) inhibited EC migration at 500 and 200 ng/ml (30%) but not at 100 ng/ml, whereas human Fc-Endostatin inhibited equally well between 500 and 100 ng/ml (25-30%) (FIG. 5). Interestingly, slightly better inhibition of EC migration was observed at 100 ng/ml for Fc-Endostatin and 200 ng/ml for endostatin (Entremed) than higher concentrations, which suggested a

U-shaped endostatin response. The best inhibition for hP1 was at 100, 62.5, and 25 ng/ml. No significant differences in inhibition between these concentrations were observed. At higher and lower concentrations less or no inhibition was observed. At 500 ng/ml of hP1 no inhibition was observed. Thus, similar to full-length endostatin, there is a U-shaped response to hP1 inhibition of EC migration. Endostatin hP2 peptide had no effect at all even at higher concentrations. However, hP6 inhibited EC migration but at higher concentrations than hP1 and no inhibition was observed for concentrations lower than 100 ng/ml.

To determine if the Zn binding site is important for anti-endothelial cell migration activity, hP1-H1/3A was also tested. This mutant peptide did show some small inhibition at 200 and 100 ng/ml. However, this inhibition was statistically not significant. Thus, peptide hP1 could inhibit VEGF-induced EC migration at equimolar concentrations (25 ng/ml) to full-length endostatin or human endostatin (EntreMed) (200 ng/ml), whereas hP6 only inhibited at doses of 100 and 200 ng/ml. Interestingly, hP1 was more potent in inhibiting EC migration than full-length endostatin. These results show that the N-terminal P1 peptide of endostatin maintains the ability to inhibit VEGF-induced EC migration and that the Zn binding site is critical for this activity.

Anti-Permeability Activities of Endostatin Peptides

The ability of endostatin peptides to inhibit VEGF-induced permeability was also tested using the Miles assay (Miles and Miles (1952) *J Physiol*, 118: 228-257). Previously, endostatin has been shown to inhibit VEGF-induced permeability using the Miles assay. Immuno-compromised SCID mice were treated 5 days before performing the Miles assay. Specifically, SCID mice were injected subcutaneously (s.c.; twice a day) with human endostatin (EntreMed; EM-ES) at a dose of 100 mg/kg/d; murine Fc-Endostatin at a dose of 20 mg/kg/d; murine endostatin peptides, mP1 and mP1-H1/3A, at a dose of 2.8 mg/kg/d or 14 mg/kg/d; or with PBS (n=3) for 5 days. At the high dose (14 mg/kg/d), both mP1 and mP1-H1/3A inhibited VEGF-induced permeability as well as human endostatin (EntreMed) and murine Fc-Endostatin (FIG. 6). However, similar results were obtained when equimolar concentrations (2.8 mg/kg/d) were used (FIG. 6). Because mP1-H1/3A showed the same inhibition as mP1, even at equimolar concentration, this suggests that there is a separation of activity between anti-tumor and anti-permeability.

Smaller peptides derived from mP1 were also shown to inhibit tumor growth. Two peptides, mP1-15 (SEQ ID NO: 118) and mP1-20 (SEQ ID NO: 108), were tested for anti-tumor activity using the LLC tumor model. Peptides were administered s.c. twice a day at dose of 2.8 mg/kg/day on days 4, 7, 10 and 14. PBS was used as a control. FIG. 7 shows that both mP1-15 and mP1-20 inhibit tumor volume. (T/C is indicated and the group size has an n equal to 5).

Thus, we have shown that a synthetic peptide, corresponding to the N-terminus of endostatin, is responsible for its anti-tumor, anti-migration, and anti-permeability activities. Zinc binding is required for the anti-tumor and anti-migration activities, because substitution of the two histidines at amino acid positions 1 and 3 in the peptide completely blocks its properties. However, Zn binding was not required for anti-permeability property.

The zinc binding requirement of endostatin for inhibiting tumor formation has been controversial, with conflicting results reported from different groups (Boehm et al. (1998) *Biochem Biophys Res Commun*, 252: 190-194; Yamaguchi et al. (1999) *Embo J*, 18: 4414-4423; Sim et al. (1999) *Angiogenesis*, 3: 41-51). Whereas, the earliest report showed that

the replacement of histidines 1 and 3 by alanines blocked the inhibitory effect of endostatin in LLC (Boehm et al. (1998) *Biochem Biophys Res Commun*, 252: 190-194), two later publications challenged this finding (Yamaguchi et al. (1999) *Embo J*, 18: 4414-4423; Sim et al. (1999) *Angiogenesis*, 3: 41-51). In one of these reports, a mutant endostatin was prepared by deleting 5 amino acids in both C— and N-termini (Yamaguchi et al. (1999) *Embo J*, 18: 4414-4423). This construct elicited anti-tumor activity, similar to full-length endostatin. However, in the employed renal Rc-9 carcinoma tumor model, the administration of endostatin was initiated when the tumor size was 300 mm³, and lasted for only 4 days, when the tumor size reached 500 mm³. The injection sites were at the periphery of the tumor, and the injection dosage was 10 µg/kg/d. In contrast, in our experiments we initiated treatment when LLC tumors reached a size of ~100 mm³ and continued until tumors were ~6000-7000 mm³. Furthermore, we treated systemically and did not inject into the periphery of the tumor.

Another publication which dealt with the relevance of zinc binding to anti-tumor activity of endostatin, demonstrated that the removal of 4 amino acids "HSHR" from N-terminus of human endostatin did not affect its anti-tumor activity (Sim et al. (1999) *Angiogenesis*, 3:41-51). Measurements of zinc binding revealed that this mutant bound 2 atoms of zinc per molecule of endostatin, whereas the wild type bound 10 atoms of zinc per endostatin molecule. However, in our crystal structure studies of endostatin, we have demonstrated that the endostatin employed for crystallization studies, contains 1 atom of zinc/endostatin molecule and the removal of the 4 amino acids "HSHR" from N-terminus lacks zinc binding (Ding et al. (1998) *Proc Natl Acad Sci USA*, 95: 10443-10448).

Endostatin is generated by proteolytic cleavage of collagen 18 (O'Reilly et al. (1997) *Cell*, 88: 277-285; Wen et al. (1999) *Cancer Res*, 59: 6052-6056; Felbor et al. (2000) *Embo J*, 19: 1187-1194). The first amino acid at the N-terminus of endostatin is a histidine. The presence of histidine is important for conferring Zn binding to endostatin. Consequently, we are led to conclude that the processing of collagen 18 to endostatin may be highly regulated.

Several groups have shown that peptides derived from endostatin have antiangiogenic effects (Wickstrom et al. (2004) *J Biol Chem*, 279: 20178-20185; Cattaneo et al. (2003) *Exp Cell Res*, 283: 230-236; Chillemi et al. (2003) *J Med Chem*, 46: 4165-4172; Morbidelli et al. (2003) *Clin Cancer Res*, 9: 5358-5369; Cho et al. (2004) *Oncol Rep*, 11: 191-195). An N-terminal peptide comprising amino acids 6-49 (lacking the zinc binding histidines) has inhibited endothelial cell proliferation and migration (Cattaneo et al. (2003) *Exp Cell Res*, 283: 230-236; Chillemi et al. (2003) *J Med Chem*, 46: 4165-4172). A Matrigel assay, employing this peptide has resulted in the inhibition of angiogenesis in vivo. However, no antitumor data was presented. In another study, a C-terminal peptide (amino acids 135-184) retaining the Cys135-Cys165 disulfide bond, has demonstrated anti-tumor activity (Morbidelli et al. (2003) *Clin Cancer Res*, 9: 5358-5369). However, the peptide was administered at the tumor periphery and not systemically. Cho et al. have shown that N-terminus, which includes the Zn binding site, and the C-terminus of endostatin are not required for anti-tumor activity (Cho et al. (2004) *Oncol Rep*, 11: 191-195). However, this peptide and full-length endostatin were not tested at equimolar concentrations. Our results differ from these groups in that the P1 peptide could inhibit tumor formation, migration, and permeability at equimolar concentrations to full length endostatin. Furthermore, at higher concentrations (14 mg/kg/

d) mP2 could inhibit LLC tumor formation as well as mP1 at 2.8 mg/kg/d. However, mP1 at 14 mg/kg/d inhibited LLC tumor formation less than at 2.8 mg/kg/d. Thus, a U-shaped curve seems to be associated with anti-tumor activity of endostatin as a function of the protein concentration. Similar results were observed for full-length endostatin using the pancreatic BxPC-3 and ASPC-1 tumor models. Therefore, determination of optimum endostatin concentration may be an important factor. In vitro assays have demonstrated a similar biphasic characteristic by endostatin such as seen in migration assays (see FIG. 5).

The fact that full-length endostatin is not required for its anti-tumor activity explains the initial inconsistencies of endostatin activity. Endostatin has two disulfide bonds. Aggregation of endostatin in *E. coli* preparations is caused by random inter-molecular disulfides after PBS dialysis. Whereas, endostatin demonstrates a single protein molecule under reducing conditions, most of the protein in an identical sample does not enter the polyacrylamide gel under nonreducing condition: It is probably the degree of nonspecific aggregation that is responsible for the lack of activity in some of the preparations. Endostatin is most likely released from the aggregate in animals over a period of time, resulting in a denatured protein or partial fragments, which are capable of demonstrating anti-tumor properties due to their N-terminal peptide. Presumably, some of the preparations yield larger aggregates, which make such a release inefficient and give rise to a product that is incapable of eliciting antiangiogenic response in mice.

What is the basis of endostatin's anti-tumor activity? A large number of mechanisms have been proposed. One which has been studied in more detail is the endostatin binding to integrin $\alpha 5 \beta 1$ (Wickstrom et al. (2002) Cancer Res, 62: 5580-5589). Based on the findings of these authors, an assembly of several cell surface proteins and components including $\alpha 5 \beta 1$, are responsible for interactions between endostatin and this integrin (Wickstrom et al. (2003) J Biol Chem, 278: 37895-37901). However, no anti-tumor data were presented to confirm the above mechanism. More recently, the same authors have shown that an 11 amino acid peptide derived from endostatin containing arginines and showing heparin binding, is responsible for antiangiogenic activity of endostatin (Wickstrom et al. (2004) J Biol Chem, 279: 20178-20185). We speculate that the phenomena observed by these investigators, reflects some of the properties associated with heparin binding characteristic of endostatin, and not its anti-tumor activity. Previously, we reported that disruption of heparin binding of endostatin (accomplished by the mutation of two discontinuous arginines on the protein surface) blocked cell motility (Kuo et al. (2001) J Cell Biol, 152: 1233-1246). Furthermore, our endostatin hP3 peptide (see Table 1), which contains the peptide reported by the authors, failed to inhibit tumor growth.

Materials and Methods:

Cell Culture and Reagents

Human BxPC-3 pancreatic adenocarcinoma and Lewis lung carcinoma (LLC) cells were grown and maintained as described earlier (Kisker et al. (2001) Cancer Res, 61:7669-7674; O'Reilly et al. (1994) Cell, 79: 315-328). For BxPC-3 tumor cell injection, cells were grown in 900-cm² roller bottles. Human microvascular endothelial cells (HMVEC-d; Clonetics, Walkersville, Md.) were cultured in microvascular endothelial cell growth medium (EGM-2 MV; Clonetics) and maintained at 5% CO₂ in a 37° C. humidified incubator. Recombinant human endostatin was a generous gift from EntreMed Corporation (Rockville, Md.) and recombinant

human and murine FcEndostatin were prepared as described earlier (Bergers et al. (1999) Science, 284: 808-812). Human and murine endostatin peptides were synthesized by SynPep Corporation (Dublin, Calif.). Peptides were resuspended in PBS or 50 mM Tris, 150 mM NaCl, pH 7.5. PECAM, purified rat anti-mouse CD31, was obtained from BD Pharmingen (San Diego, Calif.) and human recombinant VEGF was obtained from the NIH (Bethesda, Md.).

Animal Studies

All animal procedures were performed in compliance with Boston Children's Hospital guidelines, and protocols were approved by the Institutional Animal Care and Use Committee. Male (24-27 g) immunocompetent C57Bl/6J mice (Jackson Laboratories, Bar Harbor, Me.) and immunocompromised SCID mice (Massachusetts General Hospital, Boston, Mass.) were used. Mice were 7-9 weeks of age. They were acclimated, caged in groups of five in a barrier care facility, and fed with animal chow and water ad libitum. Animals were anesthetized via inhalation of isoflurane (Baxter, Deerfield, IL) before all surgical procedures and observed until fully recovered. Animals were euthanized by a lethal dose of CO₂ asphyxiation.

Tumor Models

BxPC-3 and LLC cells were grown in cell culture as described above. The cell concentration was adjusted to 50×10^6 cells/ml. Mice were shaved and the dorsal skin was cleaned with ethanol before tumor cell injection. A suspension of 5×10^6 tumor cells in 0.1 ml RPMI-1640 (for BxPC-3) or DMEM (for LLC) was injected subcutaneously (s.c.) into the dorsa of mice at the proximal midline. BxPC-3 cells were implanted in SCID mice and LLC in C57Bl/6J mice.

Animals with Lewis Lung carcinoma (600-800 mm³ tumors) were euthanized, and the skin overlying the tumor was cleaned with Betadine and ethanol. Tumor tissue was excised under aseptic conditions. A suspension of tumor cells in 0.9% normal saline was made by passage of viable tumor tissue through a sieve and a series of sequentially smaller hypodermic needles of diameter 22- to 30-gauge. The final concentration was adjusted to 1×10^7 cells/ml, and the suspension was placed on ice. The injection of tumor cells was performed as described above.

The mice were weighed and tumors were measured every 3-5 days in two diameters with a dial-caliper. Volumes were determined using the formula $a^2 \times b \times 0.52$ (where a is the shortest and b is the longest diameter). Data is represented as volume of treated tumor over control (T/C). At the completion of each experiment, the mice were euthanized with CO₂ asphyxiation. Tumors were fixed in 10% buffered Formalin (Fisher Scientific, Fair Lawn, N.J.) and paraffin-embedded.

For treatment of tumor-bearing mice, tumor volumes were allowed to grow to approximately 100 mm³, and mice were randomized. Treatment was performed by single bolus s.c. injections. Peptides were administered twice a day (every 12 hrs). Doses indicated for peptides were corrected for the purity of peptides (approximately 70%). For example, mice injected with 4 mg/kg/d peptide, were actually injected with 2.8 mg/kg/d after correction. The unpaired Student t-test was used for statistical analysis.

Immunohistochemistry

Tumors were fixed in 10% buffered Formalin overnight at 4° C. The next day, tumors were washed three times in PBS and paraffin-embedded. Sections (5 μ m) were permeabilized with 40 μ g/ml proteinase K (Roche Diagnostics Corp.) in 0.2 M Tris-HCl buffer (pH 7.6) for 25 minutes at 37° C. and washed with PBS. PECAM (1:250) was incubated at 4° C.

overnight. Stainings were amplified using tyramide signal amplification direct and indirect kits (NEN Life Science Products Inc., Boston, Mass.). Sections were photographed at 400× magnification using a NIKON TE300 microscope. Vessel density (average of three fields) was determined by IPLab software. The unpaired Student t-test was used for statistical analysis.

Cell Migration Assay

The motility response of HMVEC-d cells was assayed using a modified Boyden chamber. Cells were plated in T75-cm2 flasks at 0.5×10⁶ cells per flask and allowed to grow for 48 h (~70% confluent) prior to the migration assay. To facilitate cell adhesion, the upper membrane of a transwell (8 mm pore; Costar) was coated with fibronectin (10 mg/ml; Becton Dickinson, Bedford, Mass.) for 1 h at 37° C. Coated membranes were rinsed with PBS and allowed to air dry immediately before use. Cells were detached by trypsinization, treated with trypsinization neutralization solution (Clonetics), and resuspended at a final concentration of 1×10⁶ cells/ml in serum-free endothelial basal medium (EBM; Clonetics) containing 0.1% BSA. Cells (200,000 in 0.2 ml) were then treated with 0.2 ml of EBM/BSA containing endostatin or peptides at the indicated concentrations. Cells were incubated for 20 min. at 37° C. with occasional shaking. Cells (50,000 in 100 µl) were added to the upper chamber of the transwell. EBM or EBM supplemented with VEGF (5 ng/ml) was added to the lower chamber and cells were allowed to migrate toward the bottom chamber for 4 h in a humidified incubator containing 5% CO₂. Transwell filters were rinsed once with PBS and fixed and stained using a Diff-Quik staining kit (Baxter) following the manufacturer's protocol. Non-migrated cells were removed from the upper chamber with a cotton swab. Stained filters were cut out of the chamber and mounted onto slides using Permount (Fisher). The number of migrated cells was measured using microscopy (three fields from each membrane were captured using a 40x objective), and images were captured with a CCD camera using SPOT

software. Total migration per membrane was quantified from the captured images using Scion Image software (National Institutes of Health). All experiments were run in triplicate. The unpaired Student t-test was used for statistical analysis.

Miles Vascular Permeability Assay (The Miles Assay)

SCID mice were injected subcutaneously (s.c.) with human endostatin (EntreMed; 100 mg/kg/day), murine Fc-Endostatin (20 mg/kg/d), peptides (either 14 mg/kg/d or 2.8 mg/kg/d), and with saline (200 µl) (n=12) for 5 days prior to performing the Miles assay (25). Briefly, Evan's blue dye (100 µl of a 1% solution in PBS) was injected intravenously (i.v.) into mice. After 10 minutes, 50 µl of human recombinant VEGF (1 ng/µl) or PBS were injected intradermally into the pre-shaved back skin. After 20 minutes, the animals were euthanized and an area of skin that included the blue spot resulting from leakage of the dye was removed. Evan's blue dye was extracted from the skin by incubation with formaldehyde for 5 days at room temperature, and the absorbance of extracted dye was measured at 620 nm using a spectrophotometer. The unpaired Student t-test was used for statistical analysis.

Statistical Methods

Data are expressed as mean+S.D. Statistical significance was assessed using the Student t-test. P<0.05 was considered statistically significant.

Incorporation by Reference

The contents of all cited references (including literature references, GenBank Accession numbers, issued patents, published patent applications as cited throughout this application) are hereby expressly incorporated by reference.

Equivalents

Those skilled in the art will recognize, or be able to ascertain using no more than routine experimentation, many equivalents of the specific embodiments of the invention described herein. Such equivalents are intended to be encompassed by the following claims.

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<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 35

aacagcccc tgtcaggcgg catgcggggc atccgc 36

<210> SEQ ID NO 36
<211> LENGTH: 12
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 36

Asn Ser Pro Leu Ser Gly Gly Met Arg Gly Ile Arg
1 5 10

<210> SEQ ID NO 37
<211> LENGTH: 78
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 37

cacagccacc gcgacttcca gccggtgctc cacctggttg cgctcaacag cccctgtca 60
ggcggcatgc ggggcatac 78

<210> SEQ ID NO 38
<211> LENGTH: 26
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 38

His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Ser Pro Leu Ser Gly Gly Met Arg Gly Ile
20 25

<210> SEQ ID NO 39
<211> LENGTH: 72
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

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<400> SEQUENCE: 39
cacagccacc gcgacttcca gccggtgctc cacctggttg cgctcaacag cccctgtca 60
ggcggcatgc gg 72

<210> SEQ ID NO 40
<211> LENGTH: 24
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 40
His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Ser Pro Leu Ser Gly Gly Met Arg
20

<210> SEQ ID NO 41
<211> LENGTH: 69
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 41
cacagccacc gcgacttcca gccggtgctc cacctggttg cgctcaacag cccctgtca 60
ggcggcatg 69

<210> SEQ ID NO 42
<211> LENGTH: 23
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 42
His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Ser Pro Leu Ser Gly Gly Met
20

<210> SEQ ID NO 43
<211> LENGTH: 66
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 43
cacagccacc gcgacttcca gccggtgctc cacctggttg cgctcaacag cccctgtca 60
ggcggc 66

<210> SEQ ID NO 44
<211> LENGTH: 22
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

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<400> SEQUENCE: 44

His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Ser Pro Leu Ser Gly Gly
20

<210> SEQ ID NO 45
<211> LENGTH: 63
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 45

cacagccacc gcgacttcca gccggtgctc cacctggttg cgctcaacag cccctgtca 60
ggc 63

<210> SEQ ID NO 46
<211> LENGTH: 21
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 46

His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Ser Pro Leu Ser Gly
20

<210> SEQ ID NO 47
<211> LENGTH: 60
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 47

cacagccacc gcgacttcca gccggtgctc cacctggttg cgctcaacag cccctgtca 60

<210> SEQ ID NO 48
<211> LENGTH: 20
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 48

His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Ser Pro Leu Ser
20

<210> SEQ ID NO 49
<211> LENGTH: 57
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

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<400> SEQUENCE: 49

cacagccacc gcgacttcca gccggtgctc cacctggttg cgctcaacag cccctg 57

<210> SEQ ID NO 50
<211> LENGTH: 19
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 50

His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15

Ser Pro Leu

<210> SEQ ID NO 51
<211> LENGTH: 54
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 51

cacagccacc gcgacttcca gccggtgctc cacctggttg cgctcaacag cccc 54

<210> SEQ ID NO 52
<211> LENGTH: 18
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 52

His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15

Ser Pro

<210> SEQ ID NO 53
<211> LENGTH: 51
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 53

cacagccacc gcgacttcca gccggtgctc cacctggttg cgctcaacag c 51

<210> SEQ ID NO 54
<211> LENGTH: 17
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 54

His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15

Ser

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<210> SEQ ID NO 55
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 55

cacagccacc gcgacttcca gccggtgctc cacctggttg cgctcaac 48

<210> SEQ ID NO 56
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 56

His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15

<210> SEQ ID NO 57
<211> LENGTH: 45
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 57

cacagccacc gcgacttcca gccggtgctc cacctggttg cgctc 45

<210> SEQ ID NO 58
<211> LENGTH: 15
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 58

His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu
1 5 10 15

<210> SEQ ID NO 59
<211> LENGTH: 42
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 59

cacagccacc gcgacttcca gccggtgctc cacctggttg cg 42

<210> SEQ ID NO 60
<211> LENGTH: 14
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 60

His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala

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1	5	10	
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<div><210> SEQ ID NO 65</div> <div><211> LENGTH: 78</div> <div><212> TYPE: DNA</div> <div><213> ORGANISM: Artificial Sequence</div> <div><220> FEATURE:</div> <div><223> OTHER INFORMATION: Description of Artificial Sequence: DNA encoding exemplary anti-angiogenic peptide</div> <div><400> SEQUENCE: 65</div> <div>actcatcagg actttcagcc agtgctccac ctggtggcac tgaacacccc cctgtctgga60</div> <div>ggcatgcgtg gtatccgt78</div>			
<div><210> SEQ ID NO 66</div> <div><211> LENGTH: 26</div> <div><212> TYPE: PRT</div> <div><213> ORGANISM: Artificial Sequence</div> <div><220> FEATURE:</div> <div><223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide</div>			

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<400> SEQUENCE: 66

Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn Thr
1 5 10 15
Pro Leu Ser Gly Gly Met Arg Gly Ile Arg
20 25

<210> SEQ ID NO 67
<211> LENGTH: 75
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 67

catcaggact ttcagccagt gctccacctg gtggcactga acacccccct gtctggaggc 60
atgcgtggta tccgt 75

<210> SEQ ID NO 68
<211> LENGTH: 25
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 68

His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn Thr Pro
1 5 10 15
Leu Ser Gly Gly Met Arg Gly Ile Arg
20 25

<210> SEQ ID NO 69
<211> LENGTH: 72
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 69

caggactttc agccagtgct ccacctggtg gcactgaaca cccccctgtc tggaggcatg 60
cgtggtatcc gt 72

<210> SEQ ID NO 70
<211> LENGTH: 24
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 70

Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn Thr Pro Leu
1 5 10 15
Ser Gly Gly Met Arg Gly Ile Arg
20

<210> SEQ ID NO 71
<211> LENGTH: 69
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:

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<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 71

gactttcagc cagtgtcca cctggtggca ctgaacaccc ccctgtctgg aggcattgcgt 60

ggtatccgt 69

<210> SEQ ID NO 72
<211> LENGTH: 23
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 72

Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn Thr Pro Leu Ser
1 5 10 15

Gly Gly Met Arg Gly Ile Arg
20

<210> SEQ ID NO 73
<211> LENGTH: 67
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 73

atttcagcca gtgtccacc tgggtggcact gaacaccccc ctgtctggag gcatgcgtgg 60

tatccgt 67

<210> SEQ ID NO 74
<211> LENGTH: 22
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 74

Phe Gln Pro Val Leu His Leu Val Ala Leu Asn Thr Pro Leu Ser Gly
1 5 10 15

Gly Met Arg Gly Ile Arg
20

<210> SEQ ID NO 75
<211> LENGTH: 63
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 75

cagccagtgc tccacctggt ggactgaac acccccctgt ctggaggcat gcgtggtatc 60

cgt 63

<210> SEQ ID NO 76
<211> LENGTH: 21
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:

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<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 76

Gln Pro Val Leu His Leu Val Ala Leu Asn Thr Pro Leu Ser Gly Gly
1 5 10 15
Met Arg Gly Ile Arg
20

<210> SEQ ID NO 77
<211> LENGTH: 60
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 77

ccagtgtctcc acctgggtggc actgaacacc cccctgtctg gaggcattgcg tggatatccgt 60

<210> SEQ ID NO 78
<211> LENGTH: 20
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 78

Pro Val Leu His Leu Val Ala Leu Asn Thr Pro Leu Ser Gly Gly Met
1 5 10 15
Arg Gly Ile Arg
20

<210> SEQ ID NO 79
<211> LENGTH: 57
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 79

gtgtctccacc tgggtggcact gaacaccccc ctgtctggag gcatgcgtgg tatccgt 57

<210> SEQ ID NO 80
<211> LENGTH: 19
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 80

Val Leu His Leu Val Ala Leu Asn Thr Pro Leu Ser Gly Gly Met Arg
1 5 10 15
Gly Ile Arg

<210> SEQ ID NO 81
<211> LENGTH: 54
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA encoding exemplary anti-angiogenic peptide

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<400> SEQUENCE: 81

ctccacctgg tggcactgaa caccctcctg tctggaggca tgcgtggtat ccgt 54

<210> SEQ ID NO 82
<211> LENGTH: 18
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 82

Leu His Leu Val Ala Leu Asn Thr Pro Leu Ser Gly Gly Met Arg Gly
1 5 10 15

Ile Arg

<210> SEQ ID NO 83
<211> LENGTH: 51
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 83

cacctggtgg cactgaacac cccctgtct ggaggcatgc gtggtatccg t 51

<210> SEQ ID NO 84
<211> LENGTH: 17
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 84

His Leu Val Ala Leu Asn Thr Pro Leu Ser Gly Gly Met Arg Gly Ile
1 5 10 15

Arg

<210> SEQ ID NO 85
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 85

ctggtggcac tgaacacccc cctgtctgga ggcattcgtg gtatccgt 48

<210> SEQ ID NO 86
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 86

Leu Val Ala Leu Asn Thr Pro Leu Ser Gly Gly Met Arg Gly Ile Arg
1 5 10 15

<210> SEQ ID NO 87
<211> LENGTH: 45

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<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 87

gtggcactga acacccccct gtctggaggc atgcgtggta tccgt 45

<210> SEQ ID NO 88
<211> LENGTH: 15
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 88

Val Ala Leu Asn Thr Pro Leu Ser Gly Gly Met Arg Gly Ile Arg
1 5 10 15

<210> SEQ ID NO 89
<211> LENGTH: 42
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 89

gcactgaaca cccccctgtc tggaggcatg cgtggtatcc gt 42

<210> SEQ ID NO 90
<211> LENGTH: 14
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 90

Ala Leu Asn Thr Pro Leu Ser Gly Gly Met Arg Gly Ile Arg
1 5 10

<210> SEQ ID NO 91
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 91

ctgaacaccc ccctgtctgg aggcattcgt ggtatccgt 39

<210> SEQ ID NO 92
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 92

Leu Asn Thr Pro Leu Ser Gly Gly Met Arg Gly Ile Arg
1 5 10

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<210> SEQ ID NO 93
<211> LENGTH: 36
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 93

aacaccccc tgtctggagg catgctgggt atccgt 36

<210> SEQ ID NO 94
<211> LENGTH: 12
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 94

Asn Thr Pro Leu Ser Gly Gly Met Arg Gly Ile Arg
1 5 10

<210> SEQ ID NO 95
<211> LENGTH: 78
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 95

catactcatc aggactttca gccagtgtct cacctgggtgg cactgaacac cccctgtct 60

ggaggcatgc gtggtatc 78

<210> SEQ ID NO 96
<211> LENGTH: 26
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 96

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15

Thr Pro Leu Ser Gly Gly Met Arg Gly Ile
20 25

<210> SEQ ID NO 97
<211> LENGTH: 75
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 97

catactcatc aggactttca gccagtgtct cacctgggtgg cactgaacac cccctgtct 60

ggaggcatgc gtggt 75

<210> SEQ ID NO 98
<211> LENGTH: 25
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:

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<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 98

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Thr Pro Leu Ser Gly Gly Met Arg Gly
20 25

<210> SEQ ID NO 99
<211> LENGTH: 72
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 99

catactcatc aggactttca gccagtgtc cacctggtgg cactgaacac cccctgtct 60
ggagggcatgc gt 72

<210> SEQ ID NO 100
<211> LENGTH: 24
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 100

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Thr Pro Leu Ser Gly Gly Met Arg
20

<210> SEQ ID NO 101
<211> LENGTH: 69
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 101

catactcatc aggactttca gccagtgtc cacctggtgg cactgaacac cccctgtct 60
ggagggcatg 69

<210> SEQ ID NO 102
<211> LENGTH: 23
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 102

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Thr Pro Leu Ser Gly Gly Met
20

<210> SEQ ID NO 103
<211> LENGTH: 66
<212> TYPE: DNA

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<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 103

catactcatc aggactttca gccagtgtc cacctggtgg cactgaacac cccctgtct 60

ggaggc 66

<210> SEQ ID NO 104
<211> LENGTH: 22
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 104

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15

Thr Pro Leu Ser Gly Gly
20

<210> SEQ ID NO 105
<211> LENGTH: 63
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 105

catactcatc aggactttca gccagtgtc cacctggtgg cactgaacac cccctgtct 60

gga 63

<210> SEQ ID NO 106
<211> LENGTH: 21
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 106

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15

Thr Pro Leu Ser Gly
20

<210> SEQ ID NO 107
<211> LENGTH: 60
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 107

catactcatc aggactttca gccagtgtc cacctggtgg cactgaacac cccctgtct 60

<210> SEQ ID NO 108
<211> LENGTH: 20
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:

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<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 108

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Thr Pro Leu Ser
20

<210> SEQ ID NO 109
<211> LENGTH: 57
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 109

catactcatc aggactttca gccagtgtc cacctggtgg cactgaacac cccctg 57

<210> SEQ ID NO 110
<211> LENGTH: 19
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 110

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Thr Pro Leu

<210> SEQ ID NO 111
<211> LENGTH: 54
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 111

catactcatc aggactttca gccagtgtc cacctggtgg cactgaacac cccc 54

<210> SEQ ID NO 112
<211> LENGTH: 18
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary anti-angiogenic peptide

<400> SEQUENCE: 112

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Thr Pro

<210> SEQ ID NO 113
<211> LENGTH: 51
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 113

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catactcatc aggactttca gccagtgtc cacctggtgg cactgaacac c 51

<210> SEQ ID NO 114
<211> LENGTH: 17
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 114

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15

Thr

<210> SEQ ID NO 115
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 115

catactcatc aggactttca gccagtgtc cacctggtgg cactgaac 48

<210> SEQ ID NO 116
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 116

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15

<210> SEQ ID NO 117
<211> LENGTH: 45
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 117

catactcatc aggactttca gccagtgtc cacctggtgg cactg 45

<210> SEQ ID NO 118
<211> LENGTH: 15
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 118

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu
1 5 10 15

<210> SEQ ID NO 119
<211> LENGTH: 42
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:

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<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 119

catactcatc aggactttca gccagtgctc cacctggtgg ca 42

<210> SEQ ID NO 120
<211> LENGTH: 14
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 120

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala
1 5 10

<210> SEQ ID NO 121
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 121

catactcatc aggactttca gccagtgctc cacctggtg 39

<210> SEQ ID NO 122
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 122

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val
1 5 10

<210> SEQ ID NO 123
<211> LENGTH: 36
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 123

catactcatc aggactttca gccagtgctc cacctg 36

<210> SEQ ID NO 124
<211> LENGTH: 12
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 124

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu
1 5 10

<210> SEQ ID NO 125
<211> LENGTH: 21
<212> TYPE: PRT

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<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 125

His Ser His Arg Asp Phe Val Ala Leu Asn Ser Pro Leu Ser Gly Gly
1 5 10 15
Met Arg Gly Ile Arg
20

<210> SEQ ID NO 126
<211> LENGTH: 21
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 126

His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Leu Ser Gly Gly
1 5 10 15
Met Arg Gly Ile Arg
20

<210> SEQ ID NO 127
<211> LENGTH: 21
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 127

Gln Pro Val Leu His Leu Val Ala Leu Asn Thr Pro Leu Ser Gly Gly
1 5 10 15
Met Arg Gly Ile Arg
20

<210> SEQ ID NO 128
<211> LENGTH: 21
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 128

His Thr His Gln Asp Phe Val Ala Leu Asn Thr Pro Leu Ser Gly Gly
1 5 10 15
Met Arg Gly Ile Arg
20

<210> SEQ ID NO 129
<211> LENGTH: 21
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 129

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Leu Ser Gly Gly
1 5 10 15
Met Arg Gly Ile Arg

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20

<210> SEQ ID NO 130
<211> LENGTH: 27
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Consensus
anti-angiogenic peptide
<220> FEATURE:
<221> NAME/KEY: MOD_RES
<222> LOCATION: (2)
<223> OTHER INFORMATION: any amino acid
<220> FEATURE:
<221> NAME/KEY: MOD_RES
<222> LOCATION: (4)
<223> OTHER INFORMATION: any amino acid
<220> FEATURE:
<221> NAME/KEY: MOD_RES
<222> LOCATION: (17)
<223> OTHER INFORMATION: any amino acid

<400> SEQUENCE: 130

His Xaa His Xaa Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15

Xaa Pro Leu Ser Gly Gly Met Arg Gly Ile Arg
20 25

<210> SEQ ID NO 131
<211> LENGTH: 21
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Consensus
anti-angiogenic peptide
<220> FEATURE:
<221> NAME/KEY: MOD_RES
<222> LOCATION: (2)
<223> OTHER INFORMATION: any amino acid
<220> FEATURE:
<221> NAME/KEY: MOD_RES
<222> LOCATION: (4)
<223> OTHER INFORMATION: any amino acid
<220> FEATURE:
<221> NAME/KEY: MOD_RES
<222> LOCATION: (17)
<223> OTHER INFORMATION: any amino acid

<400> SEQUENCE: 131

His Xaa His Xaa Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15

Xaa Pro Leu Ser Gly
20

<210> SEQ ID NO 132
<211> LENGTH: 699
<212> TYPE: DNA
<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 132

gagcccaaattttgtgacaa aactcacaca tgcccaccgt gccagcacc tgaactcctg 60
gggggaccgt cagtcttctt cttcccccca aaaccaagg acacctcat gatctcccg 120
accctgagg tcacatgcgt ggtggtggac gtgagccacg aagacctga ggtcaagttc 180
aactggtacg tggacggcgt ggaggtgcat aatgccaaaga caaagccgcg ggaggagcag 240
tacaacagca cgtaccgtgt ggtcagcgtc ctcaccgtcc tgcaccagga ctggctgaat 300
ggcaaggagt acaagtgcaa ggtctccaac aaagccctcc cagcccccat cgagaaaacc 360

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atctccaaag ccaaagggca gccccgagaa ccacaggtgt acaccctgcc cccatcccgg 420
gatgagctga ccaagaacca ggtcagcctg acctgcctgg tcaaaggctt ctatcccagc 480
gacatcgccg tggagtggga gagcaatggg cagccggaga acaactacaa gaccacgcct 540
cccgtgctgg actccgacgg ctctttcttc ctctacagca agctcaccgt ggacaagagc 600
aggtggcagc aggggaacgt cttctcatgc tccgtgatgc atgaggctct gcacaaccac 660
tacacgcaga agagcctctc cctgtctcgc ggtaaataga 699

<210> SEQ ID NO 133
<211> LENGTH: 232
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 133

Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala
1 5 10 15
Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro
20 25 30
Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val
35 40 45
Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val
50 55 60
Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln
65 70 75 80
Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln
85 90 95
Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala
100 105 110
Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro
115 120 125
Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr
130 135 140
Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser
145 150 155 160
Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr
165 170 175
Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr
180 185 190
Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe
195 200 205
Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys
210 215 220
Ser Leu Ser Leu Ser Pro Gly Lys
225 230

<210> SEQ ID NO 134
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: DNA
encoding exemplary anti-angiogenic peptide

<400> SEQUENCE: 134

tctagaggtg gtctagtgcc gcgcggcagc gggtcccccg gggtgcag 48

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<210> SEQ ID NO 135
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Exemplary
anti-angiogenic peptide

<400> SEQUENCE: 135

Ser Arg Gly Gly Leu Val Pro Arg Gly Ser Gly Ser Pro Gly Leu Gln
1 5 10 15

<210> SEQ ID NO 136
<211> LENGTH: 25
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic mouse peptide

<400> SEQUENCE: 136

Met Arg Gly Ile Arg Gly Ala Asp Phe Gln Ala Phe Gln Gln Ala Arg
1 5 10 15

Ala Val Gly Leu Ser Gly Thr Phe Arg
20 25

<210> SEQ ID NO 137
<211> LENGTH: 25
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic mouse peptide

<400> SEQUENCE: 137

Thr Phe Arg Ala Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile
1 5 10 15

Val Arg Arg Ala Asp Arg Gly Ser Val
20 25

<210> SEQ ID NO 138
<211> LENGTH: 25
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic mouse peptide

<400> SEQUENCE: 138

Gly Ser Val Pro Ile Val Asn Leu Lys Asp Glu Val Leu Ser Pro Ser
1 5 10 15

Trp Asp Ser Leu Phe Ser Gly Ser Gln
20 25

<210> SEQ ID NO 139
<211> LENGTH: 25
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic mouse peptide

<400> SEQUENCE: 139

Gly Ser Gln Gly Gln Val Gln Pro Gly Ala Arg Ile Phe Ser Phe Asp
1 5 10 15

-continued

Gly Arg Asp Val Leu Arg His Pro Ala
20 25

<210> SEQ ID NO 140
<211> LENGTH: 24
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic mouse peptide

<400> SEQUENCE: 140

His Pro Ala Trp Pro Gln Lys Ser Val Trp His Gly Ser Asp Pro Ser
1 5 10 15

Gly Arg Arg Leu Met Glu Ser Tyr
20

<210> SEQ ID NO 141
<211> LENGTH: 25
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic mouse peptide

<400> SEQUENCE: 141

Glu Thr Trp Arg Thr Glu Thr Thr Gly Ala Thr Gly Gln Ala Ser Ser
1 5 10 15

Leu Leu Ser Gly Arg Leu Leu Glu Gln
20 25

<210> SEQ ID NO 142
<211> LENGTH: 26
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic mouse peptide

<400> SEQUENCE: 142

Lys Ala Ala Ser Ala His Asn Ser Tyr Ile Val Leu Ala Ile Glu Asn
1 5 10 15

Ser Phe Met Thr Ser Phe Ser Lys Lys Lys
20 25

<210> SEQ ID NO 143
<211> LENGTH: 25
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic human peptide

<400> SEQUENCE: 143

Met Arg Gly Ile Arg Gly Ala Asp Phe Gln Ala Phe Gln Gln Ala Arg
1 5 10 15

Ala Val Gly Leu Ala Gly Thr Phe Arg
20 25

<210> SEQ ID NO 144
<211> LENGTH: 25
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic human peptide

-continued

<400> SEQUENCE: 144

Thr Phe Arg Ala Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile
1 5 10 15
Val Arg Arg Ala Asp Arg Ala Ala Val
20 25

<210> SEQ ID NO 145
<211> LENGTH: 25
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic human peptide

<400> SEQUENCE: 145

Ala Ala Val Pro Ile Val Asn Leu Lys Asp Glu Leu Leu Phe Pro Ser
1 5 10 15
Trp Glu Ala Leu Phe Ser Gly Ser Glu
20 25

<210> SEQ ID NO 146
<211> LENGTH: 25
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic human peptide

<400> SEQUENCE: 146

Gly Ser Glu Gly Pro Leu Lys Pro Gly Ala Arg Ile Phe Ser Phe Asp
1 5 10 15
Gly Lys Asp Val Leu Arg His Pro Thr
20 25

<210> SEQ ID NO 147
<211> LENGTH: 24
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic human peptide

<400> SEQUENCE: 147

His Pro Thr Trp Pro Gln Lys Ser Val Trp His Gly Ser Asp Pro Asn
1 5 10 15
Gly Arg Arg Leu Thr Glu Ser Tyr
20

<210> SEQ ID NO 148
<211> LENGTH: 25
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic human peptide

<400> SEQUENCE: 148

Glu Thr Trp Arg Thr Glu Ala Pro Ser Ala Thr Gly Gln Ala Ser Ser
1 5 10 15
Leu Leu Gly Gly Arg Leu Leu Gly Gln
20 25

<210> SEQ ID NO 149

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<211> LENGTH: 28
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic human peptide

<400> SEQUENCE: 149

Leu Gly Gln Ser Ala Ala Ser Ala His His Ala Tyr Ile Val Leu Ala
1 5 10 15
Ile Glu Asn Ser Phe Met Thr Ala Ser Lys Lys Lys
20 25

<210> SEQ ID NO 150
<211> LENGTH: 27
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthesized
anti-angiogenic mutant mouse peptide

<400> SEQUENCE: 150

Ala Thr Ala Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Thr Pro Leu Ser Gly Gly Met Arg Gly Ile Arg
20 25

<210> SEQ ID NO 151
<211> LENGTH: 552
<212> TYPE: DNA
<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 151

cacagccacc gcgacttcca gccggtgctc cacctggttg cgctcaacag cccctgtca 60
ggcggcatgc ggggcatccg cggggccgac ttccagtgtt tccagcaggc gggggccgtg 120
gggctggcgg gcaccttccg cgccttcttg tctctcgccc tgcaggacct gtacagcatc 180
gtgcgccgtg ccgaccgccc agccgtgccc atcgtcaacc tcaaggacga gctgctgttt 240
cccagctggg aggtctctgtt ctcaggctct gaggggccgc tgaagcccgg ggcacgcatc 300
ttctccttta acggcaagga cgtcctgacc caccacacct ggccccagaa gagcgtgtgg 360
catggctcgg accccaacgg gcgcaggctg accgagagct actgtgagac gtggcggacg 420
gaggctccct cggccacggg ccaggcctac tcgctgtgtg ggggcaggct cctggggcag 480
agtgccgga gctgccatca cgcctacatc gtgctatgca ttgagaacag cttcatgact 540
gcctccaagt ag 552

<210> SEQ ID NO 152
<211> LENGTH: 183
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 152

His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Ser Pro Leu Ser Gly Gly Met Arg Gly Ile Arg Gly Ala Asp Phe Gln
20 25 30
Cys Phe Gln Gln Ala Arg Ala Val Gly Leu Ala Gly Thr Phe Arg Ala
35 40 45
Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala
50 55 60

-continued

Asp Arg Ala Ala Val Pro Ile Val Asn Leu Lys Asp Glu Leu Leu Phe
65 70 75 80
Pro Ser Trp Glu Ala Leu Phe Ser Gly Ser Glu Gly Pro Leu Lys Pro
85 90 95
Gly Ala Arg Ile Phe Ser Phe Asn Gly Lys Asp Val Leu Thr His Pro
100 105 110
Thr Trp Pro Gln Lys Ser Val Trp His Gly Ser Asp Pro Asn Gly Arg
115 120 125
Arg Leu Thr Glu Ser Tyr Cys Glu Thr Trp Arg Thr Glu Ala Pro Ser
130 135 140
Ala Thr Gly Gln Ala Tyr Ser Leu Leu Gly Gly Arg Leu Leu Gly Gln
145 150 155 160
Ser Ala Ala Ser Cys His His Ala Tyr Ile Val Leu Cys Ile Glu Asn
165 170 175
Ser Phe Met Thr Ala Ser Lys
180

<210> SEQ ID NO 153
<211> LENGTH: 552
<212> TYPE: DNA
<213> ORGANISM: Mus musculus

<400> SEQUENCE: 153

cataactcatc aggactttca gccagtgtct cacctggtgg cactgaacac cccctgtct 60
ggaggcatgc gtggtatccg tggagcagat ttccagtgt tccagcaagc ccgagccgtg 120
gggctgtcgg gcaccttcg ggttttctg tctctagge tgcaggatct ctatagcatc 180
gtgcgccgtg ctgaccgggg gtctgtgccc atcgtcaacc tgaaggacga ggtgctatct 240
cccagctggg actccctggt ttctggctcc caggggtcaac tgcaaccgg ggcccgcac 300
ttttcttttg acggcagaga tgtcctgaga ccccagcct ggccgcagaa gagcgtatgg 360
cacggctcgg accccagtgg gcggaggctg atggagagtt actgtgagac atggcgaact 420
gaaactactg gggctacagg tcaggcctcc tcctgtgtgt caggcaggct cctggaacag 480
aaagctgcga gctgccacaa cagctacatc gtcctgtgca ttgagaatag cttcatgacc 540
tctttctcca aa 552

<210> SEQ ID NO 154
<211> LENGTH: 184
<212> TYPE: PRT
<213> ORGANISM: Mus musculus

<400> SEQUENCE: 154

His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
1 5 10 15
Thr Pro Leu Ser Gly Gly Met Arg Gly Ile Arg Gly Ala Asp Phe Gln
20 25 30
Cys Phe Gln Gln Ala Arg Ala Val Gly Leu Ser Gly Thr Phe Arg Ala
35 40 45
Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala
50 55 60
Asp Arg Gly Ser Val Pro Ile Val Asn Leu Lys Asp Glu Val Leu Ser
65 70 75 80
Pro Ser Trp Asp Ser Leu Phe Ser Gly Ser Gln Gly Gln Leu Gln Pro
85 90 95

-continued

Gly	Ala	Arg	Ile	Phe	Ser	Phe	Asp	Gly	Arg	Asp	Val	Leu	Arg	His	Pro
			100					105					110		
Ala	Trp	Pro	Gln	Lys	Ser	Val	Trp	His	Gly	Ser	Asp	Pro	Ser	Gly	Arg
		115					120					125			
Arg	Leu	Met	Glu	Ser	Tyr	Cys	Glu	Thr	Trp	Arg	Thr	Glu	Thr	Thr	Gly
	130					135				140					
Ala	Thr	Gly	Gln	Ala	Ser	Ser	Leu	Leu	Ser	Gly	Arg	Leu	Leu	Glu	Gln
145					150					155					160
Lys	Ala	Ala	Ser	Cys	His	Asn	Ser	Tyr	Ile	Val	Leu	Cys	Ile	Glu	Asn
				165					170					175	
Ser	Phe	Met	Thr	Ser	Phe	Ser	Lys								
							180								

- The invention claimed is:
1. An isolated peptide consisting of 10 to 35 amino acids of the N-terminal region of a human endostatin protein including the first histidine of the human endostatin protein excluding SEQ ID NOs 48, 58 and 118, wherein the isolated peptide has anti-tumor or anti-angiogenic activity.
 2. The isolated peptide of claim 1, consisting of 20 to 30 amino acids.
 3. The isolated peptide of claim 1, which is selected from the group consisting of SEQ ID NOs: 2, 6, 38, 40, 42, 44, 46, 50, 52, 54, 56, 60, 62, 64, 125, 126, 130, and 131.
 4. The isolated peptide of claim 1, consisting of SEQ ID NO: 2.
 5. The isolated peptide of claim 1, consisting of SEQ ID NO: 6.
 6. The isolated peptide of claim 1, consisting of SEQ ID NOs: 64.
 7. A pharmaceutical composition comprising an isolated peptide of claim 1.
 8. A pharmaceutical composition of claim 7, further comprising a second peptide, wherein the second peptide consists of 10 to 35 amino acids of the N-terminal region of an endostatin protein including the first histidine of the endostatin protein, wherein the second peptide has anti-tumor or anti-angiogenic activity.
 9. A pharmaceutical composition of claim 7, further comprising an effective amount of zinc.
 10. An isolated peptide consisting of an amino acid sequence that is at least 95% identical to SEQ ID NO: 2, 6, 38, 40, 42, 44, 46, 50, 52, 54, 56, 60, 62, 64, 125 or 126.
 11. The isolated peptide of claim 1, wherein the peptide is linked to a heterologous peptide.
 12. The isolated peptide of claim 3, wherein the peptide is linked to a heterologous peptide.
 13. An isolated peptide consisting of 10 to 35 amino acids of the N-terminal region of an endostatin protein including the first histidine of the endostatin protein, wherein the isolated peptide has anti-tumor or anti-angiogenic activity and is not SEQ ID NO: 48, 58, 102, 108, or 118.
 14. The isolated peptide of claim 13, consisting of 20 to 30 amino acids.
 15. An isolated peptide consisting of an amino acid sequence that is at least 95% identical to SEQ ID NO: 4, 96, 98, 100, 104, 106, 110, 112, 114, 116, 120, 122, 124, 128 or 129.
 16. The isolated peptide of claim 13, which is selected from the group consisting of SEQ ID NOs: 4, 96, 98, 100, 104, 106, 110, 112, 114, 116, 120, 122, 124, 128 and 129.
 17. The isolated peptide of claim 13, consisting of SEQ ID NO: 4.
 18. The isolated peptide of claim 13, consisting of SEQ ID NO: 124.
 19. A pharmaceutical composition comprising an isolated peptide of claim 13.
 20. A pharmaceutical composition of claim 19, further comprising a second peptide, wherein the second peptide consists of 10 to 35 amino acids of the N-terminal region of an endostatin protein including the first histidine of the endostatin protein, wherein the peptide has anti-tumor or anti-angiogenic activity.
 21. A pharmaceutical composition of claim 19, further comprising an effective amount of zinc.
 22. A composition of claim 13, wherein the peptide is linked to a heterologous peptide.

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